805/2825 FIGURE 711

806/2825 FIGURE 712

MAATFFGEVVKAPCRAGTEDEEEEEEGRRETPEDREVRLQLARKREVRLLRRQTKTSLEVSLLEKYPCSKFIIAI GNNAVAFLSSFVMNSGVWEEVGCAKLWNEWCRTTDTTHLSSTEAFCVFYHLKSNPSVFLCQCSCYVAEDQQYQWL EKVFGSCPRKNMQITILTCRHVTDYKTSESTGSLPSPFLRALKTQNFKDSACCPLLEQPNIVHDLPAAVLSYCQV WKIPAILYLCYTDVMKLDLITVEAFKPILSTRSLKGLVKNIPQSTEILKKLMTTNEIQSNIYT

807/2825 FIGURE 713

ACTCAGTGTTCGCGGGAGCCGCACCTACACCAGCCAACCCAGATCCCGAGGTCCGACAGCGCCCGGCCCAGATCC CCACGCCTGCCAGGAGCAAGCCGAGAGCCAGCCGGCCGCCACTCCGACTCCGAGCAGTCTCTGTCCTTCGACC CGAGCCCGCGCCCTTTCCGGGACCCCTGCCCGCGGCCAGCGTGCCAACCTGCCGGCCATGGAGACCCCGTCC CAGCGCGCGCCACCCGCAGCGGGGCGCAGCCCAGCTCCACTCGCTGTCGCCCACCCGCATCACCCGGCTGCAG AACGCAGGGCTGCGCCTTCGCATCACCGAGTCTGAAGAGGTGGTCAGCCGCGAGGTGTCCGGCATCAAGGCCGCC $\tt CTGAGCAAAGTGCGTGAGGAGTTTAAGGAGCTGAAAGCGCGCAATACCAAGAAGGAGGGTGACCTGATAGCTGCT$ CAGGCTCGGCTGAAGGACCTGGAGGCTCTGCTGAACTCCAAGGAGGCCGCACTGAGCACTGCTCTCAGTGAGAAG CGCACGCTGGAGGCCGAGCTGCATGATCTGCGGGGCCAGGTGGCCAAGCTTGAGGCAGCCCTAGGTGAGGCCAAG AAGCAACTTCAGGATGAGATGCTGCGGCGGGTGGATGCTGAGAACAGGCTGCAGACCATGAAGGAACTGGAC TTCCAGAAGAACATCTACAGTGAGGAGCTGCGTGAGACCAAGCGCCGTCATGAGACCCGACTGGTGGAGATTGAC AATGGGAAGCAGCGTGAGTTTGAGAGCCGGCTGGCGGATGCGCTGCAGGAACTGCGGGCCCAGCATGAGGACCAG GTGGAGCAGTATAAGAAGGAGCTGGAGAAGACTTATTCTGCCAAGCTGGACAATGCCAGGCAGTCTGCTGAGAGG AACAGCAACCTGGTGGGGGCTGCCCACGAGGAGCTGCAGCAGTCGCGCATCGACAGCCTCTCTGCCCAG CTCAGCCAGCTCCAGAAGCAGCTGGCAGCCAAGGAGGCGAAGCTTCGAGACCTGGAGGACTCACTGGCCCGTGAG CGGGACACCAGCCGGCGGCTGCTGGCGGAAAAGGAGCGGGAGATGGCCGAGATGCGGGCAAGGATGCAGCAGCAG $\tt CTGGACGAGTACCAGGAGCTTCTGGACATCAAGCTGGCCCTGGACATGGAGATCCACGCCTACCGCAAGCTCTTG$ GAGGGCGAGGAGGAGGCTACCCTGTCCCCCAGCCCTACCTCGCAGCCGTGGCCGTGGCCGTGCTTCCTCAC TCATCCCAGACACAGGGTGGGGGCAGCGTCACCAAAAAGCGCAAACTGGAGTCCACTGAGAGCCGCAGCAGCTTC TCACAGCACGCACTAGCGGGCGCGTGGCCGTGGAGGAGGTGGATGAGGAGGGCAAGTTTGTCCGGCTGCGC AACAAGTCCAATGAGGACCAGTCCATGGGCAATTGGCAGATCAAGCGCCAGAATGGAGATGATCCCTTGCTGACT TACCGGTTCCCACCAAAGTTCACCCTGAAGGCTGGGCAGGTGGTGACGATCTGGGCTGCAGGAGCTGGGGCCACC CACAGCCCCCTACCGACCTGGTGTGGAAGGCACAGAACACCTGGGGCTGCGGGAACAGCCTGCGTACGGCTCTC ATCAACTCCACTGGGGAAGAAGTGGCCATGCGCAAGCTGGTGCGCTCAGTGACTGTGGTTGAGGACGACGAGGAT GAGGATGAGATGACCTGCTCCATCACCACCATGTGAGTGGTAGCCGCCGCTGAGCCGAGCCTGCACTGGGGCC ACCCAGCCAGGCCTGGGGGCAGCCTCTCCCCAGCCTCCCCGTGCCAAAAATCTTTTCATTAAAGAATGTTTTGGAA CTTT

808/2825 FIGURE **714**

METPSQRRATRSGAQASSTPLSPTRITRLQEKEDLQELNDRLAVYIDRVRSLETENAGLRLRITESEEVVSREVS GIKAAYEAELGDARKTLDSVAKERARLQLELSKVREEFKELKARNTKKEGDLIAAQARLKDLEALLNSKEAALST ALSEKRTLEGELHDLRGQVAKLEAALGEAKKQLQDEMLRRVDAENRLQTMKEELDFQKNIYSEELRETKRRHETR LVEIDNGKQREFESRLADALQELRAQHEDQVEQYKKELEKTYSAKLDNARQSAERNSNLVGAAHEELQQSRIRID SLSAQLSQLQKQLAAKEAKLRDLEDSLARERDTSRRLLAEKEREMAEMRARMQQQLDEYQELLDIKLALDMEIHA YRKLLEGEEERLRLSPSPTSQRSRGRASSHSSQTQGGGSVTKKRKLESTESRSSFSQHARTSGRVAVEEVDEEGK FVRLRNKSNEDQSMGNWQIKRQNGDDPLLTYRFPPKFTLKAGQVVTIWAAGAGATHSPPTDLVWKAQNTWGCGNS LRTALINSTGEEVAMRKLVRSVTVVEDDEDEDGDDLLHHHHVSGSRR

809/2825 FIGURE **715**

CCAAGCCCATGAGGGCCGCGCCGCCGCCGGCGGTGCTGACGAGACGGAGCTCCTGGCCCCCGAGGAGGAGCAGA GGATCA**ATG**CGGTTCAAGAATCGATTCCAGCGGTTCATGAACCATCGAGCTCCAGCCAATGGCCGCTACAAGCCA ACTTGCTATGAACATGCTGCTAACTGTTACACACACGCATTCCTCATTGTTCCGGCCATCGTGGGCAGTGCCCTC CTCCATCGGCTGTCTGATGACTGCTGGGAAAAGATAACAGCATGGATTTATGGAATGGGACTCTGTGCCCTCTTC ATCGCTTCTACAGTATTTCACATTGTATCATGGAAAAAGAGCCACTTAAGGACAGCGGAGCATTGTTTTCACATG TGTGATAGAATGGTTATCTATTTCTTCATTGCTGCTTCTTATGCTCCATGGTTAAATCTTCGTGAACTTGGACCC CTGGCATCTCATATGCGTTGGTTTATCTGGCTCATGGCAGCTGGAGGAACCATTTATGTATTTCTCTACCATGAA AAATATAAGGTGGTTGAACTCTTTTTCTATCTCACAATGGGATTCTCTCCAGCCTTGGTGGTGACATCAATGAAC AACACCGATGGACTTCAGGAACTTGCCTGTGGGGGGCTTAATTTATTGCTTGGGAGTTGTGTTCTTCAAGAGTGAT GGCATCATTCCATTTGCCCACGCCATCTGGCACCTGTTTGTGGCCACGGCAGCTGCAGTGCATTACTACGCCATT ${\tt ACTGACTTTATATCTTTTGAATATAATTACTGTGAAAGTATAAAGGCTGTGTTCTGGAATTTTCTGCCTCACAGC}$ GATGTTTACAAACCAGACTTTTATATTTTAATTTTACAGATTTTACTACATGATTTTTCTAAATTACTATGTCAG GTTGTAAAAGTCAGTGCAATAACAAACCTTCCTTTTTAAGAAGAAAATTGTTTCTATTACTTTCCCATTCACTAG ${\tt GTAAAGAATCATGGACAGAACTTACACTACTTTTTACCATGTTTCATCTTGGCATAACATGGTTCTTTTTAAAT}$ ATGGAGAAATTGCCAATTTTTAAAAACTGCAGTTTTCCAAACTTTTCTGCCAACCTCTTACTCTGAATTCAGTGC TGCTTTGGGACATATACTTGACCTAGCTTGGTTTACCAGTGATGGAAAAGTATTTTGATATCATTAACTTTTTCA AAAGATCCAACTTTTCTCTATGCCTTTGCCACATTCTCTTCAGGGTCTCTTTCCACAGCGGATAAATGTTTTTT CTGTATTATGACAGTATTGTTGTGATGGCCATCTGCTGGAAACTCCTGAAGAGCATTATGTATTACAGTGAGCAG TTTTTCCTAGAAAAATATTTAAAGCATTGTTTGACAGGTAGAAACTCATGTATCTGTAGTCCATGAGTTATATC CTGGCTCAGTGGAGTGATATTTATGTATTATTTTTACTTTTCTCTCAGTGTCTTATATTAAGATTAACATGTTGT TAATAGTTGCTTTGTTGATTAATCTCTCTTGTTGGTGTTTTAATAAATGAAATAGGCTTGCCTTTAGATCGGGTG CTGATATTGCCTGTTTCCTAGTAATGGGCTGATCAAATGATCAGTGGAATTCTTGGTTTGATGATAACCTTATTA ATTGAAATTTTTTACTGATGTGGCTTTAAAAGAGGTTTATTTTGTATATGTTTAGAACTCTCTGATTTTGATGAA TTATATGGGAGTGAGAAACAGAAGAAGTGGTATTTGCTGGCGAGTTAAATAGGCAAGGTACCCAGTGATAACACC

810/2825 FIGURE 716

MRFKNRFQRFMNHRAPANGRYKPTCYEHAANCYTHAFLIVPAIVGSALLHRLSDDCWEKITAWIYGMGLCALFIA STVFHIVSWKKSHLRTAEHCFHMCDRMVIYFFIAASYAPWLNLRELGPLASHMRWFIWLMAAGGTIYVFLYHEKY KVVELFFYLTMGFSPALVVTSMNNTDGLQELACGGLIYCLGVVFFKSDGIIPFAHAIWHLFVATAAAVHYYAIWK YLYRSPTDFMRHL

811/2825 FIGURE 717

CTGTCGCCTGCACGGTGACCCGCGTGTGCGAGGCCTTCATGGCCAGGATCCGGGTGGAGAGGCGCTGCTGACAC
CGCCGCCACACCGCCGCCACACCGCCGCTGCCTCAGTCATGCCGAAGCACGAGTTCTCTGTGGACATGACCTGTG
GAGGCTGTGCTGAAGCTGTCTCTCGGGTCCTCAATAAGCTTGGAGGAGTTAAGTATGACATTGACCTGCCCAACA
AGAAGGTCTGCATTGAATCTGAGCACAGCATGGACACTCTGCTTGCAACCCTGAAGAAAACAGGAAAGACTGTTT
CCTACCTTGGCCTTGAGTAGCAGGGGGCCTGGTCCCCACAGCCCACAGGATGGACCAAAGGGGGCAGGATGCTGAT
CCTCCCGCTGGCTTCCAGACAGACCTGGGACTTGGCAGTCATGCCGGGTGATCGTGTTCCTGCGGAGACCCTCAG
TTGTCCTATTCCTTCCTAGCTTCCCTGCAATAAATCAAGCTGCTTTTGTTGG

812/2825 FIGURE 718

MPKHEFSVDMTCGGCAEAVSRVLNKLGGVKYDIDLPNKKVCIESEHSMDTLLATLKKTGKTVSYLGLE

813/2825 FIGURE 719

GTCTTGACGTGAGCTAGCTGGCATGGCGGCCTGCATTGCAGCGGGGCACTGGGCTGCA**ATG**GGCCTAGGCCGGAG TTTCCAAGCCGCCAGGACTCTGCTCCCCCCGCCGGCCTCTATCGCCTGCAGGGTCCACGCGGGGCCTGTCCGGCA GCAGAGCACTGGGCCTTCCGAGCCCGGTGCGTTCCAACCGCCGCAAACCGGTCATCGTGGACAAGCACCGCCC AATAGAAAGAAAGATATGTTAGAAAGGAGAAAAGTACTCCACATTCCAGAGTTCTATGTTGGAAGTATTCTTCG TGTTACTACAGCTGACCCATATGCCAGTGGAAAAATCAGCCAGTTTCTGGGGATTTGCATTCAGAGATCAGGAAG AGGACTTGGAGCTACTTTCATCCTTAGGAATGTTATCGAAGGACAAGGTGTCGAGATTTGCTTTGAACTTTATAA TCCTCGGGTCCAGGAGATTCAGGTGGTCAAATTAGAGAAACGGCTGGATGATAGCTTGCTATACTTACGAGATGC CCTTCCTGAATATAGCACTTTTGATGTGAATATGAAGCCAGTAGTACAAGAGCCTAACCAAAAAGTTCCTGTTAA TGAGCTGAAAGTAAAAATGAAGCCTAAGCCCTGGTCTAAACGCTGGGAACGTCCAAATTTTAATATTAAAGGAAT CAGATTTGATCTTTGTTTAACTGAACAGCAAATGAAAGAAGCTCAGAAGTGGAATCAGCCATGGCTTGAATTTGA ATTCTGAGAATGAATTTGGTTAGTTGCAGAAGATACATTGGCTCTAAGAGGATATATTTTGAGACCAATTTAATT TTGTACACCAGTTTATTACTCTAAAAAGAGAATTACACATGCCAAATGGACCAATGTCCATTTGCTTATTGGAGG CAAAGCTACAATAGAAGTCAGAGCATCACCAGAATGGTCTTTAATGAGCATGGAACCTGAGCAAAGGGAATAGGT GGGATGAATTTTTTTTTAATTGTGAAACAATTCATAAGCACAATATGATTTACAGAATAATAAACATTCATGTA CCCACTATCAGGTTAAGAAATAGAACATTTATTAATATGTAGGAATGTTAAGAAATAAAACATTTAATAAGA

814/2825 FIGURE 720

MGLGRSFQAARTLLPPPASIACRVHAGPVRQQSTGPSEPGAFQPPPKPVIVDKHRPVEPERRFLSPEFIPRRGRT DPLKFQIERKDMLERRKVLHIPEFYVGSILRVTTADPYASGKISQFLGICIQRSGRGLGATFILRNVIEGQGVEI CFELYNPRVQEIQVVKLEKRLDDSLLYLRDALPEYSTFDVNMKPVVQEPNQKVPVNELKVKMKPKPWSKRWERPN FNIKGIRFDLCLTEQQMKEAQKWNQPWLEFDMMREYDTSKIEAAIWKEIEASKRS

(

816/2825 FIGURE 722

 ${\tt MALWRAYQRALAAHPWKVQVLTAGSLMGLGDIISQQLVERRGLQEHQRGRTLTMVSLGCGFVGPVVGGWYKVLDR} \\ {\tt FIPGTTKVDALKKMLLDQGGFAPCFLGCFLPLVGALNGLSAQDNWAKLQRDYPDALITNYYLWPAVQLANFYLVPLHYRLAVVQCVAVIWNSYLSWKAHRL} \\ {\tt CAMPAGE CONTROL OF CONTRO$

817/2825 FIGURE **723A**

AAGTGACCCAGGGAGACAAACACTTGGAGATACTTGGGGCTGAGTTTGAGCAAGACTCCCTAACCTGTGTCTGGA CAAGTCTGATGTCCTGTGTGGCCCAAGAAGAACTGACCCCGTGTCTGGAGCTCCCACCGTTATTGCATCCCTGCT GTGGCTCACCTGCTGCTGTCTCCAGGAGCCCCTGAGAAGATTTGCCTCCTCTCCCCTGCTAAGCTCCAGGTCCTG AGATTGAATTAGGGGCTGGAGCTCACTGCACTCCAGCAGTCATGGGATAGGGCCAGCGGGTGAGGTAC CGGGGCTGCTGGCCTCCTTCAACGAGGTGGGGAACACAGCCTTGATTGTGTTTTGTGAGCTATTTTTGGCAGCCGGG TGCACCGACCCCGAATGATTGGCTATGGGGCTATCCTTGTGGCCCTGGCGGGCCTGCTCATGACTCTCCCGCACT TCATCTCGGAGCCATACCGCTACGACAACACCAGCCCTGAGGATATGCCACAGGACTTCAAGGCTTCCCTGTGCC TGCCCACAACCTCGGCCCCAGCCTCGGCCCCCTCCAATGGCAACTGCTCAAGCTACACAGAAACCCAGCATCTGA GTGTGGTGGGGATCATGTTCGTGGCACAGACCCTGCTGGGCGTGGCGGGGGTGCCCATTCAGCCCTTTGGCATCT CCTACATCGATGACTTTGCCCACAACAGCAACTCGCCCCTCTACCTCGGGATCCTGTTTGCAGTGACCATGATGG GGCCAGGCCTGGCCTTTGGGCTGGGCAGCCTCATGCTGCGCCTTTATGTGGACATTAACCAGATGCCAGAAGGTG GTATCAGCCTGACCATAAAGGACCCCCGATGGGTGGGTGCCTGGTGGCTGGGTTTCCTCATCGCTGCCGGTGCAG TGGCCCTGGCTGCCATCCCCTACTTCTTCTCCCCAAGGAAATGCCCAAGGAAAAACGTGAGCTTCAGTTTCGGC GAAAGGTCTTAGCAGTCACAGACTCACCTGCCAGGAAGGGCAAGGACTCTCCCTCTAAGCAGAGCCCTGGGGAGT CCACGAAGAAGCAGGATGGCCTAGTCCAGATTGCACCAAACCTGACTGTGATCCAGTTCATTAAAGTCTTCCCCA GGGTGCTGCTGCAGACCCTACGCCACCCCATCTTCCTGCTGGTGGTCCTGTCCCAGGTATGCTTGTCATCCATGG CTGCGGGCATGGCCACCTTCCTGCCCAAGTTCCTGGAGCGCCAGTTTTCCATCACAGCCTCCTACGCCAACCTGC TGGGCCCTGTGGGATGCGGTGCCCTTTGCCTGCGGGATGCTGCTGTGCCTCTTCTTCAGCCTGCCGCTCTTCT TTATCGGCTGCTCCAGCCACCAGATTGCGGGCATCACACACCAGACCAGTGCCCACCCTGGGCTGGAGCTGTCTC CAAGCTGCATGGAGGCCTGCTCCTGCCCATTGGACGGCTTTAACCCTGTCTGCGACCCCAGCACTCGTGTGGAAT ACATCACACCCTGCCACGCAGGCTGCTCAAGCTGGGTGGTCCAGGATGCTCTGGACAACAGCCAGGTTTTCTACA CCAACTGCAGCTGCGTGGAGGGCAACCCCGTGCTGGCAGGATCCTGCGACTCAACGTGCAGCCATCTGGTGG TGCCCTTCCTGCTCCTGGTCAGCCTGGGCTCGGCCTGTCTCACCCCACACCCTCCTTCATGCTCATCC TAAGAGGAGTGAAGAAAGAAGACATTTGGCTGTGGGCATCCAGTTCATGTTCCTGAGGATTTTTGGCCTGGA TGCCCAGCCCGTGATCCACGGCAGCGCCATCGACACCACCTGTGTGCACTGGGCCCTGAGCTGTGGGCGTCGAG CTGTCTGTCGCTACTACAATAATGACCTGCTCCGAAACCGGTTCATCGGCCTCCAGTTCTTCTTCAAAACAGGTT CTGTGATCTGCTTCGCCTTAGTTTTGGCTGTCCTGAGGCAGGACAAAGAGGCCAAGGACCAAAGAGAGCAGAT CCAGCCCTGCCGTAGAGCAGCAATTGCTAGTGTCGGGGCCAGGGAAGAAGCCAGAGGATTCCCGAGTG**TGA**GCTG TCTTGGGGCCCCACCTGGCCAAGAGTAGCAGCCACAGCAGTACCTCCTCTGAGTCCTTTGCCCAAGATTGGGTGT CAAGAGCCCTGTGTTCCATTCTGGCTCCTCCACTAAATTGCTGTGTGACTTCAGGCAAGACATTGATCCTCTCTC AGCCTTTGCTTGCTAGTCTGAACCAAAGAGTTGTTTGGGCATTTGCTGTGTTGGCCATTTCTGGAGCAAGAGGGT CTTCTTCCTCCTCCCCAGCCAGCCAGCTGTCCTGGGGCCAGGCTTTCCTGGGTGGAAAGAAGTATACCTTTCC $\tt CTGGGGCCCTAGGATAGCAAAGTGAGCCATAGTGGGCCAGGCTGCCCTCCATGCTGGGCCCCAGCCCAGGTCTGC$ ACTCGCCTGGATCACCTTCTTTGAGCCTTAGCCATCTCCTGTCAGGTAGGAATGAACTTGCCAGCCTTCAGGCTC GTTCAGCTATGACCATCTGTGCGGTCAGGGTACACTCAGCTCTCCCCCAACTCCAGCAGCCTTTAAGAAGTGT TCAGGAGTGGGACACCCAGACTTGGCAGGGCCTTCAAGAGGCCTGTGTGGGGGGCCCCAGGAATCCTTAGCTGAAG CGGGGAGACTCACTCTCCATCTCAGGAAATTCTAGCCCTTGCCCTCAGGGAGCCACGGTTGAGGGTGAGGCCCAA GACCAGGGGCCACCAAAGTCCCTGTGACCTTTGTGACTAGGATCCTAATTTCTCTATTTTCTCCTGGGTGCCTGG GTCTGTGTCACCTGGGGCAGTGTGGATAATGTTTAGTTCTGTGACACTGTTTTTTTGGGGGTGGCACCTGGTTCTC CGATGCCTGGGCTGGTCAGGCCCAGGACTGTAGTGCTGGGAGCAGTAAAGCTCAGCTCTGTGTAATGAGTGAT GCTATGGCTTGCTCGTGTCTTATGATCCAATCCTTTTCTACATCAGCCCTTGTTTTGTTTTATGGCTAGTCTTAT

818/2825 FIGURE 723B

819/2825 FIGURE **724**

MGPRIGPAGEVPQVPDKETKATMGTENTPGGKASPDPQDVRPSVFHNIKLFVLCHSLLQLAQLMISGYLKSSIST VEKRFGLSSQTSGLLASFNEVGNTALIVFVSYFGSRVHRPRMIGYGAILVALAGLLMTLPHFISEPYRYDNTSPE DMPQDFKASLCLPTTSAPASAPSNGNCSSYTETQHLSVVGIMFVAQTLLGVGGVPIQPFGISYIDDFAHNSNSPL YLGILFAVTMMGPGLAFGLGSLMLRLYVDINQMPEGGISLTIKDPRWVGAWWLGFLIAAGAVALAAIPYFFFPKE MPKEKRELQFRRKVLAVTDSPARKGKDSPSKQSPGESTKKQDGLVQIAPNLTVIQFIKVFPRVLLQTLRHPIFLL VVLSQVCLSSMAAGMATFLPKFLERQFSITASYANLLIGCLSFPSVIVGIVVGGVLVKRLHLGPVGCGALCLLGM LLCLFFSLPLFFIGCSSHQIAGITHQTSAHPGLELSPSCMEACSCPLDGFNPVCDPSTRVEYITPCHAGCSSWVV QDALDNSQVFYTNCSCVVEGNPVLAGSCDSTCSHLVVPFLLLVSLGSALACLTHTPSFMLILRGVKKEDKTLAVG IQFMFLRILAWMPSPVIHGSAIDTTCVHWALSCGRRAVCRYYNNDLLRNRFIGLQFFFKTGSVICFALVLAVLRQ QDKEARTKESRSSPAVEQQLLVSGPGKKPEDSRV

820/2825 FIGURE **725A**

GTAATTGCGAGCGAGAGTGAGTGGGGCCGGGACCCGCAGAGCCGACCCTTCTCTCCCGGGCTGCGGCAGG GCAGGGCGGGGACCTCCGCGCACCAACAGAGCCGGTTCTCAGGGCGCTTTGCTCCTTGTTTTTTCCCCGGTTCTG TTTTCTCCCCTTCTCCGGAAGGCTTGTCAAGGGGTAGGAGAAAGAGACGCAAACACAAAAGTGGAAAACAGTTAA TGACCAGCCACGGCGTCCCTGCTGTGAGCTCTGGCCGCTGCCTTCCAGGGCTCCCGAGCCACACGCTGGGGGTGC TGGCTGAGGGAACATGGCTTGTTGGCCTCAGCTGAGGTTGCTGCTGTGGAAGAACCTCACTTTCAGAAGAAGACA AACATGTCAGCTGCTGGAAGTGGCCTGGCCTCTATTTATCTTCCTGATCCTGATCTCTGTTCGGCTGAGCTA CCCACCCTATGAACAACATGAATGCCATTTTCCAAATAAAGCCATCCCTGCAGGAACACTTCCTTGGGTTCA GGGGATTATCTGTAATGCCAACACCCCTGTTTCCGTTACCCGACTCCTGGGGAGGCTCCCGGAGTTGTTGGAAA CTTTAACAAATCCATTGTGGCTCGCCTGTTCTCAGATGCTCGGAGGCTTCTTTTATACAGCCAGAAAGACACCAG CATGAAGGACATGCGCAAAGTTCTGAGAACATTACAGCAGATCAAGAAATCCAGCTCAAACTTGAAGCTTCAAGA GATGCTGAGGGCTGATGTCATTCTCCACAAGGTATTTTTGCAAGGCTACCAGTTACATTTGACAAGTCTGTGCAA TGGATCAAAATCAGAAGAGATGATTCAACTTGGTGACCAAGAAGTTTCTGAGCTTTGTGGCCTACCAAGGGAGAA ACTGGCTGCAGCAGAGCGAGTACTTCGTTCCAACATGGACATCCTGAAGCCCAATCCTGAGAACACTAAACTCTAC ATCTCCCTTCCCGAGCAAGGAGCTGGCTGAAGCCACAAAAACATTGCTGCATAGTCTTGGGACTCTGGCCCAGGA GCTGTTCAGCATGAGAAGCTGGAGTGACATGCGACAGGAGGTGATGTTTCTGACCAATGTGAACAGCTCCAGCTC TCTCAACTGGTATGAGGACAACAACTACAAAGCCCTCTTTGGAGGCAATGGCACTGAGGAAGATGCTGAAACCTT CTATGACAACTCTACAACTCCTTACTGCAATGATTTGATGAAGAATTTGGAGTCTAGTCCTCTTTCCCGCATTAT CTGGAAAGCTCTGAAGCCGCTGCTCGTTGGGAAGATCCTGTATACACCTGACACTCCAGCCACAAGGCAGGTCAT GGCTGAGGTGAACAAGACCTTCCAGGAACTGGCTGTGTTCCATGATCTGGAAGGCATGTGGGAGGAACTCAGCCC CAAGATCTGGACCTTCATGGAGAACAGCCAAGAAATGGACCTTGTCCGGATGCTGTTGGACAGCAGGGACAATGA CCACTTTTGGGAACAGCAGTTGGATGGCTTAGATTGGACAGCCCAAGACATCGTGGCGTTTTTTGGCCAAGCACCC AGAGGATGTCCAGTCCAGTAATGGTTCTGTGTACACCTGGAGAGAGCTTTCAACGAGACTAACCAGGCAATCCG GACCATATCTCGCTTCATGGAGTGTGTCAACCTGAACAAGCTAGAACCCATAGCAACAGAAGTCTGGCTCATCAA CAAGTCCATGGAGCTGCTGGATGAGGGAAGTTCTGGGCTGGTATTGTGTTCACTGGAATTACTCCAGGCAGCAT TGAGCTGCCCCATCATGTCAAGTACAAGATCCGAATGGACATTGACAATGTGGAGAGGACAAATAAAATCAAGGA GCAGGATGTGGTGGAGCAGCAATCATCAGGGTGCTGACGGGCACCGAGAAGAAAACTGGTGTCTATATGCAACA GATGCCCTATCCCTGTTACGTTGATGACATCTTTCTGCGGGTGATGAGCCGGTCAATGCCCCTCTTCATGACGCT GGCCTGGATTTACTCAGTGGCTGTGATCATCAAGGGCATCGTGTATGAGAAGGAGGCACGGCTGAAAGAGACCAT GCGGATCATGGGCCTGGACAACAGCATCCTCTGGTTTAGCTGGTTCATTAGTAGCCTCATTCCTCTTCTTGTGAG CGCTGGCCTGCTAGTGGTCATCCTGAAGTTAGGAAACCTGCTGCCCTACAGTGATCCCAGCGTGGTGTTTGTCTT CCTGTCCGTGTTTGCTGTGGTGACAATCCTGCAGTGCTTCCTGATTAGCACACTCTTCTCCAGAGCCAACCTGGC GGGCTTCACACTCAAGATCTTCGCTAGCCTGCTGTCTCCTGTGGCTTTTGGGTTTGGCTGTGAGTACTTTGCCCT TTTTGAGGAGCAGGGCATTGGAGTGCAGTGGGACAACCTGTTTGAGAGTCCTGTGGAGGAAGATGGCTTCAATCT CACCACTTCGGTCTCCATGATGCTGTTTGACACCTTCCTCTATGGGGTGATGACCTGGTACATTGAGGCTGTCTT TCCAGGCCAGTACGGAATTCCCAGGCCCTGGTATTTTCCTTGCACCAAGTCCTACTGGTTTGGCGAGGAAAGTGA GCTGGGCGTGTCCATTCAGAACCTGGTAAAAGTCTACCGAGATGGGATGAAGGTGGCTGTCGATGGCCTGGCACT GAATTTTTATGAGGGCCAGATCACCTCCTTCCTGGGCCACAATGGAGCGGGGAAGACGACCACCATGTCAATCCT GACCGGGTTGTTCCCCCCGACCTCGGGCACCGCCTACATCCTGGGAAAAGACATTCGCTCTGAGATGAGCACCAT CCGGCAGAACCTGGGGGTCTGTCCCCAGCATAACGTGCTGTTTGACATGCTGACTGTCGAAGAACACATCTGGTT CTATGCCCGCTTGAAAGGGCTCTCTGAGAAGCACGTGAAGGCGGAGATGGAGCAGATGGCCCTGGATGTTGGTTT GCCATCAAGCAAGCTGAAAAGCAAAACAAGCCAGCTGTCAGGTGGAATGCAGAGAAAGCTATCTGTGGCCTTGGC CTTTGTCGGGGGATCTAAGGTTGTCATTCTGGATGAACCCACAGCTGGTGTGGACCCTTACTCCCGCAGGGGAAT ATGGGAGCTGCTGCAAATACCGACAAGGCCGCACCATTATTCTCTCTACACACCACATGGATGAAGCGGACGT

821/2825 FIGURE **725**B

TAGCACTGTGTCATACCTGAAAAAGGAGGACAGTGTTTCTCAGAGCAGTTCTGATGCTGGCCTGGGCAGCGACCA TGAGAGTGACACGCTGACCATCGATGTCTCTGCTATCTCCAACCTCATCAGGAAGCATGTGTCTGAAGCCCGGCT CTTTCATGAGATTGATGACCGGCTCTCAGACCTGGGCATTTCTAGTTATGGCATCTCAGAGACGACCCTGGAAGA AATATTCCTCAAGGTGGCCGAAGAGAGTGGGGTGGATGCTGAGACCTCAGATGGTACCTTGCCAGCAAGACGAAA CAGGCGGCCTTCGGGGACAAGCAGAGCTGTCTTCGCCCGTTCACTGAAGATGATGCTGCTGATCCAAATGATTC-TGACATAGACCCAGAATCCAGAGAGACAGACTTGCTCAGTGGGATGGCAAAGGGTCCTACCAGGTGAAAGG CTGGAAACTTACACAGCAACAGTTTGTGGCCCTTTTGTGGAAGAGACTGCTAATTGCCAGACGGAGTCGGAAAGG ATTITTTGCTCAGATTGTCTTGCCAGCTGTTTTTGTCTGCATTGCCCTTGTGTTCAGCCTGATCGTGCCACCCTT TGGCAAGTACCCCAGCCTGGAACTTCAGCCCTGGATGTACAACGAACAGTACACATTTGTCAGCAATGATGCTCC TGAGGACACGGGAACCCTGGAACTCTTAAACGCCCTCACCAAAGACCCTGGCTTCGGGACCCGCTGTATGGAAGG AAACCCAATCCCAGACACGCCCTGCCAGGCAGGGGAGGAAGAGTGGACCACTGCCCCAGTTCCCCAGACCATCAT GGACCTCTTCCAGAATGGGAACTGGACAATGCAGAACCCTTCACCTGCATGCCAGTGTAGCAGCGACAAAATCAA GAAGATGCTGCCTGTGTGTCCCCCAGGGGGCAGGGGGGCTGCCTCCTCCACAAAGAAAACAAAACACTGCAGATAT CCTTCAGGACCTGACAGGAAGAAACATTTCGGATTATCTGGTGAAGACGTATGTGCAGATCATAGCCAAAAGCTT AAAGAACAAGATCTGGGTGAATGAGTTTAGGTATGGCGGCTTTTCCCTGGGTGTCAGTAATACTCAAGCACTTCC TCCGAGTCAAGAAGTTAATGATGCCATCAAACAAATGAAGAAACACCTAAAGCTGGCCAAGGACAGTTCTGCAGA TCGATTTCTCAACAGCTTGGGAAGATTTATGACAGGACTGGACACCAAAAATAATGTCAAGGTGTGGTTCAATAA CAAGGGCTGGCATGCAATCAGCTCTTTCCTGAATGTCATCAACAATGCCATTCTCCGGGCCAACCTGCAAAAGGG AGAGAACCCTAGCCATTATGGAATTACTGCTTTCAATCATCCCCTGAATCTCACCAAGCAGCAGCTCTCAGAGGT $\verb|CTTTGTCGTATTCCTGATCCAGGAGCGGGTCAGCAAAGCAAAACACCTGCAGTTCATCAGTGGAGTGAAGCCTGT| \\$ CATCTACTGGCTCTCTAATTTTGTCTGGGATATGTGCAATTACGTTGTCCCTGCCACACTGGTCATTATCATCTT CATCTGCTTCCAGCAGAAGTCCTATGTGTCCTCCACCAATCTGCCTGTGCTAGCCCTTCTACTTTTGCTGTATGG GTGGTCAATCACACCTCTCATGTACCCAGCCTCCTTTGTGTTCAAGATCCCCAGCACAGCCTATGTGGTGCTCAC CAGCGTGAACCTCTTCATTGGCATTAATGGCAGCGTGGCCACCTTTGTGCTGGAGCTGTTCACCGACAATAAGCT GAATAATATCAATGATATCCTGAAGTCCGTGTTCTTGATCTTCCCACATTTTTGCCTGGGACGAGGGCTCATCGA CATGGTGAAAAACCAGGCAATGGCTGATGCCCTGGAAAGGTTTGGGGAGAATCGCTTTGTGTCACCATTATCTTG GGACTTGGTGGGAAACCTCTTCGCCATGGCCGTGGAAGGGGTGGTGTTCTTCCTCATTACTGTTCTGATCCA GTACAGATTCTTCATCAGGCCCAGACCTGTAAATGCAAAGCTATCTCCTCTGAATGATGAAGATGAAGATGTGAG GCGGGAAAGACAGAGAATTCTTGATGGTGGAGGCCAGAATGACATCTTAGAAATCAAGGAGTTGACGAAGATATA TAGAAGGAAGCCGGAAGCCTGCTGTTGACAGGATTTGCGTGGGCATTCCTCCTGGTGAGTGCTTTGGGCTCCTGGG AGTTAATGGGGCTGGAAAATCATCAACTTTCAAGATGTTAACAGGAGATACCACTGTTACCAGAGGAGATGCTTT CCTTAACAAAAATAGTATCTTATCAAACATCCATGAAGTACATCAGAACATGGGCTACTGCCCTCAGTTTGATGC TGGCAAGGTTGGTGAGTGGGCGATTCGGAAACTGGGCCTCGTGAAGTATGGAGAAAAATATGCTGGTAACTATAG TGGAGGCAACAACGCAAGCTCTCTACAGCCATGGCTTTGATCGGCGGGCCTCCTGTGGTGTTTCTGGATGAACC CACCACAGGCATGGATCCCAAAGCCCGGCGGTTCTTGTGGAATTGTGCCCTAAGTGTTGTCAAGGAGGGGAGATC AGTAGTGCTTACATCTCATAGTATGGAAGAATGTGAAGCTCTTTGCACTAGGATGGCAATCATGGTCAATGGAAG GTTCAGGTGCCTTGGCAGTGTCCAGCATCTAAAAAATAGGTTTGGAGATGGTTATACAATAGTTGTACGAATAGC AGGGTCCAACCCGGACCTGAAGCCTGTCCAGGATTTCTTTGGACTTGCATTTCCTGGAAGTGTTCTAAAAGAGAA ACACCGGAACATGCTACAATACCAGCTTCCATCTTCATTATCTTCTCTGGCCAGGATATTCAGCATCCTCTCCCA GAGCAAAAAGCGACTCCACATAGAAGACTACTCTGTTTCTCAGACAACACTTGACCAAGTATTTGTGAACTTTGC CAAGGACCAAAGTGATGACCACTTAAAAGACCTCTCATTACACAAAAACCAGACAGTAGTGGACGTTGCAGT AAAGTAAAGAGGAACTAGACTTTCCTTTGCACCATGTGAAGTGTTGTGGGAGAAAAGAGCCAGAAGTTGATGTGGG

822/2825 FIGURE **725**C

AAGAAGTAAACTGGATACTGTACTGATACTATTCAATGCAATGCAATTCAATGCAATGAAAACAAAATTCCATTA CAGGGGCAGTGCCTTTGTAGCCTATGTCTTGTATGGCTCTCAAGTGAAAGACTTGAATTTAGTTTTTTACCTATA TGTGTATTCTCATTGGGGTTGCAACAATAATTCATCAAGTAATCATGGCCAGCGATTATTGATCAAAATCAAAAG GTAATGCACATCCTCATTCACTAAGCCATGCCATGCCCAGGAGACTGGTTTCCCGGTGACACATCCATTGCTGGC AATGAGTGTGCCAGAGTTATTAGTGCCAAGTTTTTCAGAAAGTTTGAAGCACCATGGTGTCATGCTCACTTTT GTGAAAGCTGCTCTGCTCAGAGTCTATCAACATTGAATATCAGTTGACAGAATGGTGCCATGCGTGGCTAACATC CTGCTTTGATTCCCTCTGATAAGCTGTTCTGGTGGCAGTAACATGCAACAAAAATGTGGGTGTCTCCAGGCACGG GAAACTTGGTTCCATTGTTATATTGTCCTATGCTTCGAGCCATGGGTCTACAGGGTCATCCTTATGAGACTCTTA AATATACTTAGATCCTGGTAAGAGGCAAAGAATCAACAGCCAAACTGCTGGGGCTGCAAGCTGCTGAAGCCAGGG CATGGGATTAAAGAGATTGTGCGTTCAAACCTAGGGAAGCCTGTGCCCATTTGTCCTGACTGTCTGCTAACATGG TACACTGCATCTCAAGATGTTTATCTGACACAAGTGTATTATTTCTGGCTTTTTGAATTAATCTAGAAAATGAAA AGATGGAGTTGTATTTTGACAAAAATGTTTGTACTTTTTAATGTTATTTGGAATTTTAAGTTCTATCAGTGACTT $\tt CTGAATCCTTAGAATGGCCTCTTTGTAGAACCCTGTGGTATAGAGGAGTATGGCCACTGCCCCACTATTTTTATT$ TTCTTATGTAAGTTTGCATATCAGTCATGACTAGTGCCTAGAAAGCAATGTGATGGTCAGGATCTCATGACATTA TGTAGCTGAAAGAGTATGTACGTACGTATAAGACTAGAGAGATATTAAGTCTCAGTACACTTCCTGTGCCATGTT ATTCAGCTCACTGGTTTACAAATATAGGTTGTCTTGTGGTTGTAGGAGCCCACTGTAACAATACTGGGCAGCCTT TTTTTTTTTTTTTTAATTGCAACAATGCAAAAGCCAAGAAAGTATAAGGGTCACAAGTCTAAACAATGAATTCT TCAACAGGGAAAACAGCTAGCTTGAAAACTTGCTGAAAAACACAACTTGTGTTTAATGGCATTTAGTACCTTCAAA TAATTGGCTTTGCAGATATTGGATACCCCATTAAATCTGACAGTCTCAAATTTTTCATCTCTTCAATCACTAGTC AAGAAAATATAAAAACAACAAATACTTCCATATGGAGCATTTTTCAGAGTTTTCTAACCCAGTCTTATTTTTCT AGAGAACTATTGTTTGGGGAAGTTCAAGTGATCTTTCAATATCATTACTAACTTCTTCCACTTTTTCCAGAATTT GAATATTAACGCTAAAGGTGTAAGACTTCAGATTTCAAATTAATCTTTCTATATTTTTTAAATTTACAGAATATT ATATAACCCACTGCTGAAAAAGAAAAAATGATTGTTTTAGAAGTTAAAGTCAATATTGATTTTAAATATAAGTA ATGAAGGCATATTTCCAATAACTAGTGATATGGCATCGTTGCATTTTACAGTATCTTCAAAAATACAGAATTTAT AGAATAATTTCTCCTCATTTAATATTTTTCAAAATCAAAGTTATGGTTTCCTCATTTTACTAAAATCGTATTCTA ATTCTTCATTATAGTAAATCTATGAGCAACTCCTTACTTCGGTTCCTCTGATTTCAAGGCCATATTTTAAAAAAT CAAAAGGCACTGTGAACTATTTTGAAGAAAACACAACATTTTAATACAGATTGAAAGGACCTCTTCTGAAGCTAG AAACAATCTATAGTTATACATCTTCATTAATACTGTGTTACCTTTTAAAATAGTAATTTTTTACATTTTCCTGTG TGTACCTATGTGAGTTTCAGAAATTCTCAAAATACGTGTTCAAAAATTTCTGCTTTTTGCATCTTTTGGGACACCTC AGAAAACTTATTAACAACTGTGAATATGAGAAATACAGAAGAAAATAATAAGCCCTCTATACATAAATGCCCAGC ACAATTCATTGTTAAAAAAACAACCAAACCTCACACTACTGTATTTCATTATCTGTACTGAAAGCAAATGCTTTGT GACTATTAAATGTTGCACATCATTCATTCACTGTATAGTAATCATTGACTAAAGCCATTTGTCTGTGTTTTTCTTC TTGTGGTTGTATATATCAGGTAAAATATTTTCCAAAGAGCCATGTGTCATGTAATACTGAACCACTTTGATATTG AGACATTAATTTGTACCCTTGTTATTATCTACTAGTAATAATGTAATACTGTAGAAATATTGCTCTAATTCTTTT CAAAATTGTTGCATCCCCTTAGAATGTTTCTATTTCCATAAGGATTTAGGTATGCTATTATCCCTTCTTATACC GATCTATTTTTGCACTGGAATATCTGAGAATTGCAAAACTAGACAAAAGTTTCACAACAGATTTCTAAGTTAAAT CATTTTCATTAAAAGGAAAAAAAAAATTTTGTATGTCAATAACTTTATATGAAGTATTAAAATGCATATT TCTATGTTGTAATATAATGAGTCACAAAATAAAGCTGTGACAGTTCTGTTGGTCTACAGAAA

823/2825 FIGURE 726

MPSAGTLPWVQGIICNANNPCFRYPTPGEAPGVVGNFNKSIVARLFSDARRLLLYSQKDTSMKDMRKVLRTLQQI $\tt KKSSSNLKLQDFLVDNETFSGFLYHNLSLPKSTVDKMLRADVILHKVFLQGYQLHLTSLCNGSKSEEMIQLGDQE$ VSELCGLPREKLAAAERVLRSNMDILKPILRTLNSTSPFPSKELAEATKTLLHSLGTLAQELFSMRSWSDMRQEV MFLTNVNSSSSSTQIYQAVSRIVCGHPEGGGLKIKSLNWYEDNNYKALFGGNGTEEDAETFYDNSTTPYCNDLMK NLESSPLSRIIWKALKPLLVGKILYTPDTPATRQVMAEVNKTFQELAVFHDLEGMWEELSPKIWTFMENSOEMDL VRMLLDSRDNDHFWEQQLDGLDWTAQDIVAFLAKHPEDVQSSNGSVYTWREAFNETNQAIRTISRFMECVNLNKL EPIATEVWLINKSMELLDERKFWAGIVFTGITPGSIELPHHVKYKIRMDIDNVERTNKIKDGYWDPGPRADPFED MRYVWGGFAYLQDVVEQAIIRVLTGTEKKTGVYMQQMPYPCYVDDIFLRVMSRSMPLFMTLAWIYSVAVIIKGIV YEKEARLKETMRIMGLDNSILWFSWFISSLIPLLVSAGLLVVILKLGNLLPYSDPSVVFVFLSVFAVVTILQCFL ISTLFSRANLAAACGGIIYFTLYLPYVLCVAWQDYVGFTLKIFASLLSPVAFGFGCEYFALFEEQGIGVQWDNLF ESPVEEDGFNLTTSVSMMLFDTFLYGVMTWYIEAVFPGQYGIPRPWYFPCTKSYWFGEESDEKSHPGSNOKRISE ${\tt ICMEEEPTHLKLGVSIQNLVKVYRDGMKVAVDGLALNFYEGQITSFLGHNGAGKTTTMSILTGLFPPTSGTAYIL}$ GKDIRSEMSTIRQNLGVCPQHNVLFDMLTVEEHIWFYARLKGLSEKHVKAEMEQMALDVGLPSSKLKSKTSOLSG GMQRKLSVALAFVGGSKVVILDEPTAGVDPYSRRGIWELLLKYRQGRTIILSTHHMDEADVLGDRIAIISHGKLC CVGSSLFLKNQLGTGYYLTLVKKDVESSLSSCRNSSSTVSYLKKEDSVSQSSSDAGLGSDHESDTLTIDVSAISN LIRKHVSEARLVEDIGHELTYVLPYEAAKEGAFVELFHEIDDRLSDLGISSYGISETTLEEIFLKVAEESGVDAE TSDGTLPARRNRRAFGDKQSCLRPFTEDDAADPNDSDIDPESRETDLLSGMDGKGSYQVKGWKLTQQQFVALLWK RLLIARRSRKGFFAQIVLPAVFVCIALVFSLIVPPFGKYPSLELQPWMYNEQYTFVSNDAPEDTGTLELLNALTK DPGFGTRCMEGNPIPDTPCQAGEEEWTTAPVPQTIMDLFQNGNWTMQNPSPACQCSSDKIKKMLPVCPPGAGGLP PPQRKQNTADILQDLTGRNISDYLVKTYVQIIAKSLKNKIWVNEFRYGGFSLGVSNTQALPPSQEVNDATKQMKK $\verb+HLKLAKDSSADRFLNSLGRFMTGLDTRNNVKVWFNNKGWHAISSFLNVINNAILRANLQKGENPSHYGITAFNHP+ \\$ LNLTKQQLSEVAPMTTSVDVLVSICVIFAMSFVPASFVVFLIQERVSKAKHLOFISGVKPVIYWLSNFVWDMCNY VVPATLVIIIFICFQQKSYVSSTNLPVLALLLLLYGWSITPLMYPASFVFKIPSTAYVVLTSVNLFIGINGSVAT ${\tt FVLELFTDNKLNNINDILKSVFLIFPHFCLGRGLIDMVKNQAMADALERFGENRFVSPLSWDLVGRNLFAMAVEG}$ VVFFLITVLIQYRFFIRPRPVNAKLSPLNDEDEDVRRERQRILDGGGQNDILEIKELTKIYRRKRKPAVDRICVG IPPGECFGLLGVNGAGKSSTFKMLTGDTTVTRGDAFLNRNSILSNIHEVHQNMGYCPQFDAITELLTGREHVEFF ALLRGVPEKEVGKVGEWAIRKLGLVKYGEKYAGNYSGGNKRKLSTAMALI

824/2825 FIGURE **727**A

GGCCGGGACCCGCAGAGCCGACCCTTCTCTCCCGGGCTGCGGCAGGGCAGGGCGGGGAGCTCCGCGCACC AACAGAGCCGGTTCTCAGGGCGCTTTGCTCCTTGTTTTTTCCCCGGGTTCTGTTTTCTCCCCCTTCTCCGGAAGGCT TGTCAAGGGGTAGGAGAAGAGACGCAAACACAAAAGTGGAAAACAGTTAATGACCAGCCACGGGCGTCCCTGCT GGCCTCAGCTGAGGTTGCTGCTGTGGAAGAACCTCACTTTCAGAAGAAGACAACATGTCAGCTGTTACTGGAAG GCCATTTTCCAAATAAAGCCATGCCCTCTGCAGGAACACTTCCTTGGGTTCAGGGGATTATCTGTAATGCCAACA ACCCCTGTTTCCGTTACCCGACTCCTGGGGAGGCTCCCGGAGTTGTTGGAAACTTTAACAAATCCATTGTGGCTC GCCTGTTCTCAGATGCTCGGAGGCTTCTTTTATACAGCCAGAAAGACACCAGCATGAAGGACATGCGCAAAGTTC TGAGAACATTACAGCAGATCAAGAAATCCAGCTCAAACTTGAAGCTTCAAGATTTCCTGGTGGACAATGAAACCT TCTCTGGGTTCCTATATCACAACCTCTCTCCCCAAAGTCTACTGTGGACAAGATGCTGAGGGCTGATGTCATTC TCCACAAGGTATTTTTGCAAGGCTACCAGTTACATTTGACAAGTCTGTGCAATGGATCAAAATCAGAAGAGATGA TTCAACTTGGTGACCAAGAAGTTTCTGAGCTTTGTGGCCTACCAAAGGAGAAACTGGCTGCAGCAGAGCGAGTAC TTCGTTCCAACATGGACATCCTGAAGCCAATCCTGAGAACACTAAACTCTACATCTCCCTTCCCGAGCAAGGAGC TGGCTGAAGCCACAAAAACATTGCTGCATAGTCTTGGGACTCTGGCCCAGGAGCTGTTCAGCATGAGAAGCTGGA GTGACATGCGACAGGAGGTGATGTTTCTGACCAATGTGAACAGCTCCAGCTCCTCCACCCAAATCTACCAGGCTG ACTACAAAGCCCTCTTTGGAGGCAATGGCACTGAGGAAGATGCTGAAACCTTCTATGACAACTCTACAACTCCTT ACTGCAATGATTTGATGAAGAATTTGGAGTCTAGTCCTCTTTCCCGCATTATCTGGAAAGCTCTGAAGCCGCTGC TCGTTGGGAAGATCCTGTATACACCTGACACTCCAGCCACAAGGCAGGTCATGGCTGAGGTGAACAAGACCTTCC AGGAACTGGCTGTGTTCCATGATCTGGAAGGCATGTGGGAGGAACTCAGCCCCAAGATCTGGACCTTCATGGAGA ACAGCCAAGAAATGGACCTTGTCCGGATGCTGTTGGACAGCAGGGACAATGACCACTTTTGGGAACAGCAGTTGG ATGGCTTAGATTGGACAGCCCAAGACATCGTGGCGTTTTTTGGCCAAGCACCCAGAGGATGTCCAGTCCAGTAATG GTTCTGTGTACACCTGGAGAGAGCTTTCAACGAGACTAACCAGGCAATCCGGACCATATCTCGCTTCATGGAGT GTGTCAACCTGAACAAGCTAGAACCCATAGCAACAGAAGTCTGGCTCATCAACAAGTCCATGGAGCTGCTGGATG AGAGGAAGTTCTGGGCTGGTATTGTGTTCACTGGAATTACTCCAGGCAGCATTGAGCTGCCCCATCATGTCAAGT ACAAGATCCGAATGGACATTGACAATGTGGAGAGGACAAATAAAATCAAGGATGGGTACTGGGACCCTGGTCCTC TCATCAGGGTGCTGACGGGCACCGAGAAGAAAACTGGTGTCTATATGCAACAGATGCCCTATCCCTGTTACGTTG ATGACATCTTTCTGCGGGTGATGAGCCGGTCAATGCCCCTCTTCATGACGCTGGCCTGGATTTACTCAGTGGCTG TGATCATCAAGGGCATCGTGTATGAGAAGGAGGCACGGCTGAAAGAGACCATGCGGATCATGGGCCTGGACAACA GCATACTCTGGTTTAGCTGGTTCATTAGTAGCCTCATTCCTCTTCTTGTGAGCGCTGGCCTGCTAGTGGTCATCC TGAAGTTAGGAAACCTGCTGCCCTACAGTGATCCCAGCGTGGTGTTTGTCTTCCTGTCCGTGTTTGCTGTGGTGA CAATCCTGCAGTGCTTCCTGATTAGCACACTCTTCTCCAGAGCCAACCTGGCAGCAGCCTGTGGGGGGCATCATCT ACTTCACGCTGTACCTGCCCTACGTCCTGTGTGTGGCATGGCAGGACTACGTGGGCTTCACACTCAAGATCTTCG TGCAGTGGGACAACCTGTTTGAGAGTCCTGTGGAGGAAGATGGCTTCAATCTCACCACTTCGATCTCCATGATGC TGTTTGACACCTTCCTCTATGGGGTGATGACCTGGTACATTGAGGCTGTCTTTCCAGGCCAGTACGGAATTCCCA GGCCCTGGTATTTTCCTTGCACCAAGTCCTACTGGTTTGGCGAGGAAAGTGATGAGAAGAGCCACCCTGGTTCCA TGGTAAAAGTCTACCGAGATGGGATGAAGGTGGCTGTCGATGGCCTGGCACTGAATTTTTATGAGGGCCAGATCA CCTCCTTCCTGGGCCACAATGGAGCGGGGAAGACGACCACCATGTCAATCCTGACCGGGTTGTTCCCCCCGACCT CGGGCACCGCCTACATCCTGGGAAAAGACATTCGCTCTGAGATGAGCACCATCCGGCAGAACCTGGGGGTCTGTC $\verb|CCCAGCATAACGTGCTGTTTGACATGCTGACTGTCGAAGAACACATCTGGTTCTATGCCCGCTTGAAAGGGCTCT|\\$ AAACAAGCCAGCTGTCAGGTGGAATGCAGAGAAAGCTATCTGTGGCCTTTGGCCTTTGTCGGGGGGATCTAAGGTTG TCATTCTGGATGAACCCACAGCTGGTGTGGACCCTTACTCCCGCAGGGGAATATGGGAGCTGCTGAAATACC GACAAGGCCGCACCATTATTCTCTCTACACACCACATGGATGAAGCGGACGTCCTGGGGGACAGGATTGCCATCA

825/2825 FIGURE **727**B

TCTCCCATGGGAAGCTGTGTGTGTGGGCTCCTCCCTGTTTCTGAAGAACCAGCTGGGAACAGGCTACTACCTGA CCTTGGTCAAGAAGATGTGGAATCCTCCCTCAGTTCCTGCAGAAACAGTAGTAGCACTGTGTCATACCTGAAAA AGGAGGACAGTGTTTCTCAGAGCAGTTCTGATGCTGGCCTGGGCAGCGACCATGAGAGTGACACGCTGACCATCG ATGTCTCTGCTATCTCCAACCTCATCAGGAAGCATGTGTCTGAAGCCCGGCTGGTGGAAGACATAGGGCATGAGC TGACCTATGTGCTGCCATATGAAGCTGCTAAGGAGGGGGCCTTTGTGGAACTCTTTCATGAGATTGATGACCGGC TCTCAGACCTGGGCATTTCTAGTTATGGCATCTCAGAGACGACCCTGGAAGAAATATTCCTCAAGGTGGCCGAAG AGAGTGGGGTGGATGCTGAGACCTCAGATGGTACCTTGCCAGCAAGACGAAACAGGCGGGCCTTCGGGGACAAGC AGAGCTGTCTTCGCCCGTTCACTGAAGATGATGCTGCTGATCCAAATGATTCTGACATAGACCCAGAATCCAGAG AGACAGACTTGCTCAGTGGGATGGATGGCAAAGGGTCCTACCAGGTGAAAGGCTGGAAACTTACACAGCAACAGT TTGTGGCCCTTTTGTGGAAGAGACTGCTAATTGCCAGACGGAGTCGGAAAGGATTTTTTGCTCAGATTGTCTTGC CAGCTGTGTTTGTCTGCATTGCCCTTGTGTTCAGCCTGATCGTGCCACCCTTTGGCAAGTACCCCAGCCTGGAAC TTCAGCCCTGGATGTACAACGAACAGTACACATTTGTCAGCAATGATGCTCCTGAGGACACGGGAACCCTGGAAC GCCAGGCAGGGGAGGAAGAGTGGACCACTGCCCCAGTTCCCCAGACCATCATGGACCTCTTCCAGAATGGGAACT GGACAATGCAGAACCCTTCACCTGCATGCCAGTGTAGCAGCGACAAAATCAAGAAGATGCTGCCTGTGTGTCCCC CAGGGGCAGGGGGCTGCCTCCACAAAGAAAACAAAACCTGCAGATATCCTTCAGGACCTGACAGGAAGAA ACATTTCGGATTATCTGGTGAAGACGTATGTGCAGATCATAGCCAAAAGCTTAAAGAACAAGATCTGGGTGAATG AGTTTAGGTATGGCGGCTTTTCCCTGGGTGTCAGTAATACTCAAGCACTTCCTCCGAGTCAAGAAGTTAATGATG CCATCAAACAAATGAAGAAACACCTAAAGCTGGCCAAGGACAGTTCTGCAGATCGATTTCTCAACAGCTTGGGAA GATTTATGACAGGACTGGACACCAGAAATAATGTCAAGGTGTGGTTCAATAACAAGGGCTGGCATGCAATCAGCT CTTTCCTGAATGTCATCAACAATGCCATTCTCCGGGCCAACCTGCAAAAGGGAGAACCCTAGCCATTATGGAA TTACTGCTTTCAATCATCCCCTGAATCTCACCAAGCAGCAGCTCTCAGAGGTGGCTCTGATGACCACATCAGTGG AGCGGGTCAGCAAAACACCTGCAGTTCATCAGTGGAGTGAAGCCTGTCATCTACTGGCTCTCTAATTTTG TCTGGGATATGTGCAATTACGTTGTCCCTGCCACACTGGTCATTATCATCTTCATCTGCTTCCAGCAGAAGTCCT ATGTGTCCTCCACCAATCTGCCTGTGCTAGCCCTTCTACTTTTGCTGTATGGGTGGTCAATCACACCTCTCATGT ACCCAGCCTCCTTTGTGTTCAAGATCCCCAGCACAGCCTATGTGGTGCTCACCAGCGTGAACCTCTTCATTGGCA TTAATGGCAGCGTGGCCACCTTTGTGCTGGAGCTGTTCACCGACAATAAGCTGAATAATATCAATGATATCCTGA AGTCCGTGTTCTTGATCTTCCCACATTTTTGCCTGGGACGAGGGCTCATCGACATGGTGAAAAACCAGGCAATGG CTGATGCCCTGGAAAGGTTTGGGGAGAATCGCTTTGTGTCACCATTATCTTGGGACTTGGTGGGACGAAACCTCT TCGCCATGGCCGTGGAAGGGGTGTTTCTTCCTCATTACTGTTCTGATCCAGTACAGATTCTTCATCAGGCCCA GACCTGTAAATGCAAAGCTATCTCCTCTGAATGATGAAGATGAAGATGTGAGGCGGGAAAGACAGAGAATTCTTG TTGACAGGATTTGCGTGGGCATTCCTCCTGGTGAGTGCTTTGGGCTCCTGGGAGTTAATGGGGCTGGAAAATCAT CAACTTTCAAGATGTTAACAGGAGATACCACTGTTACCAGAGGAGATGCTTTCCTTAACAAAAATAGTATCTTAT CAAACATCCATGAAGTACATCAGAACATGGGCTACTGCCCTCAGTTTGATGCCATCACAGAGCTGTTGACTGGGA GAGAACACGTGGAGTTCTTTGCCCTTTTGAGAGGAGTCCCAGAGAAAAGAAGTTGGCAAGGTTGGTGAGTGGGCGA TTCGGAAACTGGGCCTCGTGAAGTATGGAGAAAAATATGCTGGTAACTATAGTGGAGGCAACAAACGCAAGCTCT CTACAGCCATGGCTTTGATCGGCGGGCCTCCTGTGGTGTTTCTGGATGAACCCACCACAGGCATGGATCCCAAAG CCCGGCGGTTCTTGTGGAATTGTGCCCTAAGTGTTGTCAAGGAGGGGAGATCAGTAGTGCTTACATCTCATAGTA TGGAAGAATGTGAAGCTCTTTGCACTAGGATGGCAATCATGGTCAATGGAAGGTTCAGGTGCCTTGGCAGTGTCC AGCATCTAAAAAATAGGTTTGGAGATGGTTATACAATAGTTGTACGAATAGCAGGGTCCAACCCGGACCTGAAGC CTGTCCAGGATTTCTTTGGACTTGCATTTCCTGGAAGTGTTCTAAAAGAGAAACACCGGAACATGCTACAATACC AGCTTCCATCTTCATTATCTTCTCTGGCCAGGATATTCAGCATCCTCTCCCAGAGCAAAAAGCGACTCCACATAG AAGACTACTCTGTTTCTCAGACAACACTTGACCAAGTATTTGTGAACTTTGCCAAGGACCAAAGTGATGACC ACTTAAAAGACCTCTCATTACACAAAAACCAGACAGTAGTGGACGTTGCAGTTCTCACATCTTTTCTACAGGATG AGAAAGTGAAAGAAAGCTATGTA<u>TGA</u>AGAATCCTGTTCATACGGGGTGGCTGAAAGTAAAGAGGAACTAGACTTT $\verb|CCTTTGCACCATGTGAAGTGTTGTGGAGAAAAGAGCCAGAAGTTGATGTGGGAAGAAGTAAACTGGATACTGTAC| \\$

826/2825 FIGURE 727C

TGATACTATTCAATGCAATTCAATGCAATGAAAACAAAATTCCATTACAGGGGCAGTGCCTTTGTAGCCT ATGTCTTGTATGGCTCTCAAGTGAAAGACTTGAATTTAGTTTTTTACCTATACCTATGTGAAACTCTATTATGGA CAATAATTCATCAAGTAATCATGGCCAGCGATTATTGATCAAAAATCAAAAGGTAATGCACATCCTCATTCACTAA GCCATGCCATGCCCAGGAGACTGGTTTCCCGGTGACACATCCATTGCTGGCAATGAGTGTGCCAGAGTTATTAGT GCCAAGTTTTCAGAAAGTTTGAAGCACCATGGTGTCATGCTCACTTTTGTGAAAGCTGCTCTGCTCAGAGTC TATCAACATTGAATATCAGTTGACAGAATGGTGCCATGCGTGGCTAACATCCTGCTTTGATTCCCTCTGATAAGC TGTTCTGGTGGCAGTAACATGCAACAAAAATGTGGGTGTCTCTAGGCACGGGAAACTTGGTTCCATTGTTATATT GTCCTATGCTTCGAGCCATGGGTCTACAGGGTCATCCTTATGAGACTCTTAAATATACTTAGATCCTGGTAAGAG GCAAAGAATCAACAGCCAAACTGCTGGGGCTGCAAGCTGCTGAAGCCAGGGCATGGGATTAAAGAGATTGTGCGT TCAAACCTAGGGAAGCCTGTGCCCATTTGTCCTGACTGTCTGCTAACATGGTACACTGCATCTCAAGATGTTTAT CTGACACAAGTGTATTATTTCTGGCTTTTTGAATTAATCTAGAAAATGAAAAGATGGAGTTGTATTTTGACAAAA ATGTTTGTACTTTTAATGTTATTTGGAATTTTAAGTTCTATCAGTGACTTCTGAATCCTTAGAATGGCCTCTTT GTAGAACCCTGTGGTATAGAGGAGTATGGCCACTGCCCCACTATTTTTATTTTCTTATGTAAGTTTGCATATCAG TAGGATACTCTTAATCTCACTTCATCAATCAAATATTTTTTTGAGTGTATGCTGTAGCTGAAAGAGTATGTACGTA CGTATAAGACTAGAGAGATATTAAGTCTCAGTACACTTCCTGTGCCATGTTATTCAGCTCACTGGTTTACAAATA AAACTTGCTGAAAAACACAACTTGTGTTTATGGCATTTAGTACCTTCAAATAATTGGCTTTGCAGATATTGGATA CCCCATTAAATCTGACAGTCTCAAATTTTTCATCTCTTCAATCACTAGTCAAGAAAAATATAAAAAACAACAAATA CTTCCATATGGAGCATTTTTCAGAGTTTTCTAACCCAGTCTTATTTTTCTAGTCAGTAAACATTTGTAAAAATAC TGTTTCACTAATACTTACTGTTAACTGTCTTGAGAGAAAAGAAAAATATGAGAGAACTATTGTTTGGGGAAGTTC AAGTGATCTTTCAATATCATTACTAACTTCTTCCACTTTTTCCAAAATTTGAATATTAACGCTAAAGGTGTAAGA CTTCAGATTTCAAATTAATCTTTCTATATTTTTTAAATTTACAGAATATTATAAACCCACTGCTGAAAAAGAAA AAAATGATTGTTTTAGAAGTTAAAGTCAATATTGATTTTAAATATAAGTAATGAAGGCATATTTCCAATAACTAG TGATATGGCATCGTTGCAATTTACAGTATCTTCAAAAATACAGAATTTATAGAATAATTTCTCCTCATTTAATAT TTTTCAAAATCAAAGTTATGGTTTCCTCATTTTACTAAAATCGTATTCTAATTCTTCATTATAGTAAATCTATGA GCAACTCCTTACTTCGGTTCCTCTGATTTCAAGGCCATATTTTAAAAAAATCAAAAGGCACTGTGAACTATTTTGA AGAAAACACGACATTTTAATACAGATTGAAAGGACCTCTTCTGAAGCTAGAAACAATCTATAGTTATACATCTTC ATTAATACTGTGTTACCTTTTAAAATAGTAATTTTTTTACATTTTTCCTGTGTAAACCTAATTGTGGTAGAAATTTT TACCAACTCTATACTCAATCAAGCAAAATTTCTGTATATTCCCTGTGGAATGTACCTATGTGAGTTTCAGAAATT CTCAAAATACGTGTTCAAAAATTTCTGCTTTTGCATCTTTTGGGACACCTCAGAAAACTTATTAACAACTGTGAAT ATGAGAAATACAGAAGAAAATAATAAGCCCTCTATACATAAATGCCCCAGCACAATTCATTGTTAAAAAACAACCA AACCTCACACTACTGTATTTCATTATCTGTACTGAAAGCAAATGCTTTGTGACTATTAAATGTTGCACATCATTC ATTCACTGTATAGTAATCATTGACTAAAGCCATTTGCTGTTTTTCTTCTTGTGGXTGXATATATCAGGTAAAAT ATTTTCCAAAGAGCCATGTGTCATGTAATACTGAACCCTTTGATATTGAGACATTAATTTGGACCCTTGGTATTA TCTACTAGAATAATGTAATACTGXAGAAATATTGCTCTAATTCTTTCAAAATGGTGCATCCCCCTTAAAAXGTTC TATTTCCATAAGGATTTAGCTTGCTTATCCCTTCTTATACCCTAAGATGAAGCTGTTTTTTGTGCTCTTTGTTCAT CATTGGCCCTCATTCCAAGCACTTTACGCTGTCTGTAATGGGATCTATTTTTTGCACTGGAATATCTGAGAATTGC TGTATGTCAATAACTTTATATGAAGTATTAAAATGCATATTTCTATGTTGTAATATAATGAGTCACAAAATAAAG AAAAAAAAAAAAAA

827/2825 FIGURE **728**

MACWPQLRLLLWKNLTFRRRQTCQLLLEVAWPLFIFLILISVRLSYPPYEQHECHFPNKAMPSAGTLPWVOGIIC ${\tt NANNPCFRYPTPGEAPGVVGNFNKSIVARLFSDARRLLLYSQKDTSMKDMRKVLRTLQQIKKSSSNLKLQDFLVD}$ NETFSGFLYHNLSLPKSTVDKMLRADVILHKVFLQGYQLHLTSLCNGSKSEEMIQLGDQEVSELCGLPKEKLAAA ERVLRSNMDILKPILRTLNSTSPFPSKELAEATKTLLHSLGTLAQELFSMRSWSDMRQEVMFLTNVNSSSSSTQI YQAVSRIVCGHPEGGGLKIKSLNWYEDNNYKALFGGNGTEEDAETFYDNSTTPYCNDLMKNLESSPLSRIIWKAL KPLLVGKILYTPDTPATRQVMAEVNKTFQELAVFHDLEGMWEELSPKIWTFMENSQEMDLVRMLLDSRDNDHFWE QQLDGLDWTAQDIVAFLAKHPEDVQSSNGSVYTWREAFNETNQAIRTISRFMECVNLNKLEPIATEVWLINKSME LLDERKFWAGIVFTGITPGSIELPHHVKYKIRMDIDNVERTNKIKDGYWDPGPRADPFEDMRYVWGGFAYLQDVV EQAIIRVLTGTEKKTGVYMQQMPYPCYVDDIFLRVMSRSMPLFMTLAWIYSVAVIIKGIVYEKEARLKETMRIMG LDNSILWFSWFISSLIPLLVSAGLLVVILKLGNLLPYSDPSVVFVFLSVFAVVTILQCFLISTLFSRANLAAACG GIIYFTLYLPYVLCVAWQDYVGFTLKIFASLLSPVAFGFGCEYFALFEEQGIGVQWDNLFESPVEEDGFNLTTSI SMMLFDTFLYGVMTWYIEAVFPGQYGIPRPWYFPCTKSYWFGEESDEKSHPGSNQKRMSEICMEEEPTHLKLGVS IQNLVKVYRDGMKVAVDGLALNFYEGQITSFLGHNGAGKTTTMSILTGLFPPTSGTAYILGKDIRSEMSTIRQNL ${\tt GVCPQHNVLFDMLTVEEHIWFYARLKGLSEKHVKAEMEQMALDVGLPSSKLKSKTSQLSGGMQRKLSVALAFVGG}$ SKVVILDEPTAGVDPYSRRGIWELLLKYRQGRTIILSTHHMDEADVLGDRIAIISHGKLCCVGSSLFLKNQLGTG YYLTLVKKDVESSLSSCRNSSSTVSYLKKEDSVSQSSSDAGLGSDHESDTLTIDVSAISNLIRKHVSEARLVEDI GHELTYVLPYEAAKEGAFVELFHEIDDRLSDLGISSYGISETTLEEIFLKVAEESGVDAETSDGTLPARRNRRAF GDKQSCLRPFTEDDAADPNDSDIDPESRETDLLSGMDGKGSYQVKGWKLTQQQFVALLWKRLLIARRSRKGFFAQ IVLPAVFVCIALVFSLIVPPFGKYPSLELQPWMYNEQYTFVSNDAPEDTGTLELLNALTKDPGFGTRCMEGNPIP DTPCQAGEEEWTTAPVPQTIMDLFQNGNWTMQNPSPACQCSSDKIKKMLPVCPPGAGGLPPPQRKQNTADILQDL TGRNISDYLVKTYVQIIAKSLKNKIWVNEFRYGGFSLGVSNTQALPPSQEVNDAIKQMKKHLKLAKDSSADRFLN SLGRFMTGLDTRNNVKVWFNNKGWHAISSFLNVINNAILRANLQKGENPSHYGITAFNHPLNLTKQQLSEVALMT TSVDVLVSICVIFAMSFVPASFVVFLIQERVSKAKHLQFISGVKPVIYWLSNFVWDMCNYVVPATLVIIIFICFQ QKSYVSSTNLPVLALLLLLYGWSITPLMYPASFVFKIPSTAYVVLTSVNLFIGINGSVATFVLELFTDNKLNNIN t DILKSVFLIFPHFCLGRGLIDMVKNQAMADALERFGENRFVSPLSWDLVGRNLFAMAVEGVVFFLITVLIQYRFFIRPRPVNAKLSPLNDEDEDVRRERQRILDGGGQNDILEIKELTKIYRRKRKPAVDRICVGIPPGECFGLLGVNGA GKSSTFKMLTGDTTVTRGDAFLNKNSILSNIHEVHQNMGYCPQFDAITELLTGREHVEFFALLRGVPEKEVGKVG EWAIRKLGLVKYGEKYAGNYSGGNKRKLSTAMALIGGPPVVFLDEPTTGMDPKARRFLWNCALSVVKEGRSVVLT SHSMEECEALCTRMAIMVNGRFRCLGSVQHLKNRFGDGYTIVVRIAGSNPDLKPVQDFFGLAFPGSVLKEKHRNM LQYQLPSSLSSLARIFSILSQSKKRLHIEDYSVSQTTLDQVFVNFAKDQSDDDHLKDLSLHKNQTVVDVAVLTSF LODEKVKESYV

828/2825 FIGURE **729A**

GCGAGCGCAGCGGAGCCTGGAGAAGAGCCGCTGGGCTGCGAGGGCCGAGGGCGCGAGGGCAACCGGA ACGCCTTGCCCGCCCAGGTGGCATTTACACCCTACGCCCCGGAGCCCGGGAGCACATGCCGGCTCAGAGAATACT ATGACCAGACAGCTCAGATGTGCTGCAGCAAATGCTCGCCGGGCCAACATGCAAAAGTCTTCTGTACCAAGACCT CGGACACCGTGTGTGACTCCTGTGAGGACAGCACATACACCCAGCTCTGGAACTGGGTTCCCGAGTGCTTGAGCT GTGGCTCCCGCTGTAGCTCTGACCAGGTGGAAACTCAAGCCTGCACTCGGGAACAGAACCGCATCTGCACCTGCA GGCCCGGCTGGTACTGCGCGCTGAGCAAGCAGGAGGGGTGCCGGCTGTGCGCCGCCGCCGCCCGG GCTTCGGCGTGGCCAGACCAGGAACTGAAACATCAGACGTGGTGTGCAAGCCCTGTGCCCCGGGGACGTTCTCCA ACACGACTTCATCCACGGATATTTGCAGGCCCCACCAGATCTGTAACGTGGTGGCCATCCCTGGGAATGCAAGCA GCCCCAGCCCCAGCTGAAGGGAGCACTGGCGACTTCGCTCTTCCAGTTGGACTGATTGTGGGTGTGACAGCCT TGGGTCTACTAATAATAGGAGTGGTGAACTGTGTCATCATGACCCAGGTGAAAAAGAAGCCCTTGTGCCTGCAGA GAGAAGCCAAGGTGCCTCACTTGCCTGCCGATAAGGCCCGGGGTACACAGGGCCCCGAGCAGCACCTGCTGA ACCAGCCACAGGCACCAGGCGTGGAGGCCAGTGGGGCCGGGGAGGCCCGGGAGCTCAGATTCTT $\verb|CCCCTGGTGGCCATGGGACCCAGGTCAATGTCACCTGCATCGTGAACGTCTGTAGCAGCTCTGACCACAGCTCAC| \\$ AGTGCTCCTCCCAAGCCAGCTCCACAATGGGAGACACAGATTCCAGCCCCTCGGAGTCCCCGAAGGACGAGCAGG TCCCCTTCTCCAAGGAGGAATGTGCCTTTCGGTCACAGCTGGAGACCCCAGAGACCCTGCTGGGGAGCACCGAAG $\tt AGAAGCCCCTGCCCCTTGGAGTGCCTGATGCTGGGATGAAGCCCAGT{\color{red}{\textbf{TAA}}} CCAGGCCGGTGTGGGCTGTGTCGTA$ GCCAAGGTGGGCTGAGCCCTGGCAGGATGACCCTGCGAAGGGGCCCTGGTCCTTCCAGGCCCCCACCACTAGGAC CCAGAAAACCCCAGCATCCTTTTCTGCAGAGGGGCTTTCTGGAGAGGAGGGATGCTGCCTGAGTCACCCATGAAG CTGTAGGGAACGGGGTCCTTCAAGTTAGCTCAGGAGGCTTGGAAAGCATCACCTCAGGCCAGGTGCAGTGGCTCA CGCCTATGATCCCAGCACTTTGGGAGGCTGAGGCGGGTGGATCACCTGAGGTTAGGAGTTCGAGACCAGCCTGGC CAACATGGTAAAACCCCATCTCTACTAAAAATACAGAAATTAGCCGGGCGTGGTGGCGGGCACCTATAGTCCCAG CTACTCAGAAGCCTGAGGCTGGGAAATCGTTTGAACCCGGGAAGCGGAGGTTGCAGGGAGCCGAGATCACGCCAC GCCAACGACAAAGCCAAACTCTGCCAGCCACATCCAACCCCCCACCTGCCATTTGCACCCTCCGCCTTCACTCCG GTGTGCCTGCAGCCCCGCGCCTCCTTCCTTGCTGTCCTAGGCCACACCATCTCCTTTCAGGGAATTTCAGGAACT AGAGATGACTGAGTCCTCGTAGCCATCTCTCTACTCCTACCTCAGCCTAGACCCTCCTCCTCCTCCCCAGAGGGGTG GGTTCCTCTCCCCACTCCCACCTTCAATTCCTGGGCCCCAAACGGGCTGCCCTGCCACTTTGGTACATGGCCA TTGAATGGCCTGCCTTGAAGCCACTGAAGCTGGGATTCCTCCCCATTAGAGTCAGCCTTCCCCCTCCCAGGGCCA GGGCCCTGCAGAGGGGAAACCAGTGTAGCCTTGCCCGGATTCTGGGAGGAAGCAGGTTGAGGGGCTCCTGGAAAG GCTCAGTCTCAGGAGCATGGGGATAAAGGAGAAGGCATGAAATTGTCTAGCAGAGCAGGGGCAGGGTGATAAATT GTTGATAAATTCCACTGGACTTGAGCTTGGCAGCTGAACTATTGGAGGGTGGGAGAGCCCAGCCATTACCATGGA GGCTGAGGGGAGGATCACTGGAGCCCAGGAGTTTGAGGCTGCAGCGAGCTATGATCGCGCCACTACACTCCAGCC CATTGGACCCACATGAGGACATGATGGAGCGCACCTGCCCCCTGGTGGACAGTCCTGGGAGAACCTCAGGCTTCC

829/2825 FIGURE **729B**

830/2825 FIGURE 730

MAPVAVWAALAVGLELWAAAHALPAQVAFTPYAPEPGSTCRLREYYDQTAQMCCSKCSPGQHAKVFCTKTSDTVC DSCEDSTYTQLWNWVPECLSCGSRCSSDQVETQACTREQNRICTCRPGWYCALSKQEGCRLCAPLRKCRPGFGVA RPGTETSDVVCKPCAPGTFSNTTSSTDICRPHQICNVVAIPGNASMDAVCTSTSPTRSMAPGAVHLPQPVSTRSQ HTQPTPEPSTAPSTSFLLPMGPSPPAEGSTGDFALPVGLIVGVTALGLLIIGVVNCVIMTQVKKKPLCLQREAKV PHLPADKARGTQGPEQQHLLITAPSSSSSSLESSASALDRRAPTRNQPQAPGVEASGAGEARASTGSSDSSPGGH GTQVNVTCIVNVCSSSDHSSQCSSQASSTMGDTDSSPSESPKDEQVPFSKEECAFRSQLETPETLLGSTEEKPLP LGVPDAGMKPS

831/2825 FIGURE **731A**

GGCGCGGCCTGGAGCCCCGGGAGCGCGCGCGCGCGCCCAGCCGGCTCTCCTGGCCTCGCGCTGCA CGCGACTCCCGTTCCTATTCACCCTGGTCGCACTGCTGCCGCCCGGAGCTCTCTGCGAAGTCTGGACGCAGAGGC TGCACGGCGCCAGCGCCCTTGCCCCAGGACCGGGGCTTCCTCGTGGTGCAGGGCGACCCGCGCGAGCTGCGGC TGTGGGCGCGGGGATGCCAGGGGGGCGAGCCGCGGAGAAGCCGCTCCGGAGGAAACGGAGCGCTGCCC TGCAGCCCGAGCCCATCAAGGTGTACGGACAGGTTAGTCTGAATGATTCCCACAATCAGATGGTGGTGCACTGGG CTGGAGAGAAAAGCAACGTGATCGTGGCCTTGGCCCGAGATAGCCTGGCATTGGCCGAGGCCCAAGAGCAGTGATG TGTACGTGTCTTACGACTATGGAAAATCATTCAAGAAAATTTCAGACAAGTTAAACTTTGGCTTGGGAAATAGGA GTGAAGCTGTTATCGCCCAGTTCTACCACAGCCCTGCGGACAACAAGCGGTACATCTTTGCAGACGCTTATGCCC AGTACCTCTGGATCACGTTTGACTTCTGCAACACTCTTCAAGGCTTTTCCATCCCATTTCGGGCAGCTGATCTCC TCCTACACAGTAAGGCCTCCAACCTTCTCTTGGGCTTTGACAGGTCCCACCCCAACAAGCAGCTGTGGAAGTCAG ATGACTTTGGCCAGACCTGGATCATGATTCAGGAACATGTCAAGTCCTTTTCTTGGGGAATTGATCCCTATGACA AACCAAATACCATCTACATTGAACGACACGAACCCTCTGGCTACTCCACTGTCTTCCGAAGTACAGATTTCTTCC AGTCCCGGGAAAACCAGGAAGTGATCCTTGAGGAAGTGAGAGATTTTCAGCTTCGGGACAAGTACATGTTTGCTA $\tt TGAGAGCCCCAGTTTGTCACAAGACATCCTATTAATGAATATTACATCGCAGATGCCTCCGAGGACCAGGTGT$ TTGTGTGTGTCAGCCACAGTAACAACCGCACCAATTTATACATCTCAGAGGCAGAGGGGCTGAAGTTCTCCCTGT $\verb|CCTTGGAGAACGTGCTCTATTACAGCCCAGGAGGGGCCGGCAGTGACACCTTGGTGAGGTATTTTGCAAATGAAC| \\$ CATTTGCTGACTTCCACCGAGTGGAAGGATTGCAAGGAGTCTACATTGCTACTCTGATTAATGGTTCTATGAATG TCAACCTCCAGCTCCGGAGAATGCCCATCCTGTCCAAGGAGTCGGCTCCAGGCCTCATCATCGCCACTGGCTCAG $\tt TGGGAAGGACTTGGCTAGCAAGACAAACGTGTACATCTCTAGCAGTGCTGGAGCCAGGTGGCGAGAGGCACTTC$ CTGGACCTCACTACACATGGGGAGACCACGGCGGAATCATCACGGCCATTGCCCAGGGCATGGAAACCAACG AGCTAAAATACAGTACCAATGAAGGGGAGACCTGGAAAACATTCATCTTCTCTGAGAAGCCAGTGTTTGTGTATG GCCTCCTCACAGAACCTGGGGAGAAGAGCACTGTCTTCACCATCTTTGGCTCGAACAAAGAGAATGTCCACAGCT GGCTGATCCTCCAGGTCAATGCCACGGATGCCTTGGGAGTTCCCTGCACAGAGAATGACTACAAGCTGTGGTCAC CATCTGATGAGCGGGGGAATGAGTGTTTGCTGGGACACAAGACTGTTTTCAAACGGCGGACCCCCATGCCACAT GCTTCAATGGAGAGGACTTTGACAGGCCGGTGGTCGTGCCAACTGCTCCTGCACCCGGGAGGACTATGAGTGTG ACTTCGGTTTCAAGATGAGTGAAGATTTGTCATTAGAGGTTTGTGTTCCAGATCCGGAATTTTCTGGAAAGTCAT ACTCCCCTCCTGTGCCTTGCCCTGTGGGTTCTACTTACAGGAGACGAGAGGCTACCGGAAGATTTCTGGGGACA CTTGTAGCGGAGGAGATGTTGAAGCGCGACTGGAAGGAGAGCTGGTCCCCTGTCCCCTGGCAGAAGAACGAGT TCATTCTGTATGCTGTGAGGAAATCCATCTACCGCTATGACCTGGCCTCGGGAGCCACCGAGCAGTTGCCTCTCA $\verb|CCGGGCTACGGGCAGCAGTGGCCCTGGACTTTGACTATGAGCACAACTGTTTGTATTGGTCCGACCTGGCCTTGG|\\$ ACGTCATCCAGCGCCTCTGTTTGAATGGAAGCACAGGGCAAGAGGTGATCATCAATTCTGGCCTGGAGACAGTAG AAGCTTTGGCTTTTGAACCCCTCAGCCAGCTGCTTTACTGGGTAGATGCAGGCTTCAAAAAGATTGAGGTAGCTA ATCCAGATGGCGACTTCCGACTCACAATCGTCAATTCCTCTGTGCTTGATCGTCCCAGGGCTCTGGTCCTCGTGC CTGCTGCCTATCACCTGGTGTCTGAGGATGTGAAGTGGCCCAATGGCATCTCTGTGGACGACCAGTGGATTTACT GGACGGATGCCTACCTGGAGTGCATAGAGCGGATCACGTTCAGTGGCCAGCAGCGCTCTGTCATTCTGGACAACC TCCCGCACCCCTATGCCATTGCTGTCTTTAAGAATGAAATCTACTGGGATGACTGGTCACAGCTCAGCATATTCC GAGCTTCCAAATACAGTGGGTCCCAGATGGAGATTCTGGCAAACCAGCTCACGGGGCTCATGGACATGAAGATTT TCTACAAGGGGAAGAACACTGGAAGCAATGCCTGTGTGCCCAGGCCATGCAGCCTGCTGTGCCCAAGGCCA ACAACAGTAGAAGCTGCAGGTGTCCAGAGGATGTGTCCAGCAGTGTGCTTCCATCAGGGGACCTGATGTGTGACT GCCCTCAGGGCTATCAGCTCAAGAACAATACCTGTGTCAAAGAAGAGAACACCTGTCTTCGCAACCAGTATCGCT GCAGCAACGGGAACTGTATCAACAGCATTTGGTGGTGTGACTTTGACAACGACTGTGGAGACATGAGCGATGAGA GAAACTGCCCTACCACCATCTGTGACCTGGACACCCAGTTTCGTTGCCAGGAGTCTGGGACTTGTATCCCACTGT CCTATAAATGTGACCTTGAGGATGACTGTGGAGACAACAGTGATGAAAGTCATTGTGAAATGCACCAGTGCCGGA

832/2825 FIGURE **731B**

GTGACGAGTACAACTGCAGTTCCGGCATGTGCATCCGCTCCTCCTGGGTATGTGACGGGGACAACGACTGCAGGG ACTGGTCTGATGAAGCCAACTGTACCGCCATCTATCACACCTGTGAGGCCTCCAACTTCCAGTGCCGAAACGGGC ACTGCATCCCCCAGCGGTGGGCGTGTGACGGGGATACGGACTGCCAGGATGGTTCCGATGAGGATCCAGTCAACT GTGAGAAGAAGTGCAATGGATTCCGCTGCCCAAACGGCACTTGCATCCCATCCAGCAAACATTGTGATGGTCTGC GTGATTGCTCTGATGGCTCCGATGAACAGCACTGCGAGCCCCTCTGTACGCACTTCATGGACTTTGTGTGTAAGA ACCGCCAGCAGTGCCTGTTCCACTCCATGGTCTGTGACGGAATCATCCAGTGCCGCGACGGGTCCGATGAGGATG CGGCGTTTGCAGGATGCTCCCAAGATCCTGAGTTCCACAAGGTATGTGATGAGTTCGGTTTCCAGTGTCAGAATG AAAACCCCACAGAAGCCCCAAACTGCTCCCGCTACTTCCAGTTTCGGTGTGAGAATGGCCACTGCATCCCCAACA GATGGAAATGTGACAGGGAGAACGACTGTGGGGACTGGTCTGATGAGAAGGATTGTGGAGATTCACATATTCTTC CCTTCTCGACTCCTGGGCCCTCCACGTGTCTGCCCAATTACTACCGCTGCAGCAGTGGGACCTGCGTGATGGACA CTGCTGCCTCCACTCCACCCAACTTGGGCGATGTGACCGATTTGAGTTCGAATGCCACCAACCGAAGACGTGTA GCACCTTGACTTGCATGAGCAGGGAGTTCCAGTGCGAGGACGGGGAGGCCTGCATTGTGCTCTCGGAGCGCTGCG ACGGCTTCCTGGACTGCTCGGACGAGAGCGATGAAAAGGCCTGCAGTGATGAGTTGACTGTACAAAGTACAGA $\verb|CTTGTGTATATATGTCTACTACAGGGTGGTTGGAGAGAGCATATGGAAGACTCTGGAGACCCACAGCAATAAGA| \\$ CAAACACTGTATTAAAAGTCTTGAAACCAGATACCACGTATCAGGTTAAAGTACAGGTTCAGTGTCTCAGCAAGG CACACAACACCAATGACTTTGTGACCCTGAGGACCCCAGAGGGATTGCCAGATGCCCCTCGAAATCTCCAGCTGT $\tt CACTCCCCAGGGAAGCAGAAGGTGTGATTGTAGGCCACTGGGCTCCTCCCATCCACACCCCATGGCCTCATCCGTG$ AGTACATTGTAGAATACAGCAGGAGTGGTTCCAAGATGTGGGCCTCCCAGAGGGCTGCTAGTAACTTTACAGAAA TCAAGAACTTATTGGTCAACACTCTATACACCGTCAGAGTGGCTGCGGTGACTAGTCGTGGAATAGGAAACTGGA GCGATTCTAAATCCATTACCACCATAAAAGGAAAAGTGATCCCACCAGATATCCACATTGACAGCTATGGTG AAAATTATCTAAGCTTCACCCTGACCATGGAGAGTGATATCAAGGTGAATGGCTATGTGGTGAACCTTTTCTGGG CATTTGACACCCACAAGCAAGAGAGAGAACTTTGAACTTCCGAGGAAGCATATTGTCACACAAAGTTGGCAATC ATGTTATGACCAGAGGGGTTCGCCCACCTGCACCTAGCCTCAAGGCCAAAGCCATCAACCAGACTGCAGTGGAAT ${\tt GTACCTGGACCGGCCCCGGAATGTGGTTTATGGTATTTTCTATGCCACGTCCTTTCTTGACCTCTATCGCAACC}$ CGAAGAGCTTGACTACTTCACTCCACAACAAGACGGTCATTGTCAGTAAGGATGAGCAGTATTTGTTTCTGGTCC GTGTAGTGGTACCCTACCAGGGGCCATCCTCTGACTACGTTGTAGTGAAGATGATCCCGGACAGCAGGCTTCCAC $\verb|CCCGTCACCTGCATGTGGTTCATACGGGCAAAACCTCCGTGGTCATCAAGTGGGAATCACCGTATGACTCTCCTG|\\$ GTAACAGCACTGTGGAATACACCCTTAACAAGTTGGAGCCTGGCGGGAAATACCACATCATTGTCCAACTGGGGA ACATGAGCAAAGATTCCAGCATAAAAATTACCACAGTTTCATTATCAGCACCTGATGCCTTAAAAAATCATAACAG AAAATGATCATGTTCTTCTGTTTTGGAAAAGCCTGGCTTTAAAGGAAAAGCATTTTAATGAAAGCAGGGGCTATG AGATACACATGTTTGATAGTGCCATGAATATCACAGCTTACCTTGGGAATACTACTGACAATTTCTTTAAAATTT AGCCTGCCATCCTGCTGTACGATGAGCTGGGGTCTGGTGCAGATGCATCTGCAACGCAGGCTGCCAGATCTACGG ATGTTGCTGCTGTGGTGGTGCCCATCTTATTCCTGATACTGCTGAGCCTGGGGGTTGGGGTTTGCCATCCTGTACA CGAAGCACCGGAGGCTGCAGAGCAGCTTCACCGCCTTCGCCAACAGCCACTACAGCTCCAGGCTGGGGTCCGCAA TCTTCTCCTCTGGGGGATGACCTGGGGGAAGATGATGAAGATGCCCCTATGATAACTGGATTTTCAGATGACGTCC CCATGGTGATAGCCTGAAAGAGCTTTCCTCACTAGAAACCAAATGGTGTAAATATTTTATTTGATAAAGATAGTT AAAAA

833/2825 FIGURE **732**

 ${\tt MATRSSRRESRLPFLFTLVALLPPGALCEVWTQRLHGGSAPLPQDRGFLVVQGDPRELRLWARGDARGASRADEK}$ $\verb"PLRRKRSAALQPEPIKVYGQVSLNDSHNQMVVHWAGEKSNVIVALARDSLALARPKSSDVYVSYDYGKSFKKISD"$ HPNKQLWKSDDFGQTWIMIQEHVKSFSWGIDPYDKPNTIYIERHEPSGYSTVFRSTDFFQSRENQEVILEEVRDF QLRDKYMFATKVVHLLGSEQQSSVQLWVSFGRKPMRAAQFVTRHPINEYYIADASEDQVFVCVSHSNNRTNLYIS EAEGLKFSLSLENVLYYSPGGAGSDTLVRYFANEPFADFHRVEGLQGVYIATLINGSMNEENMRSVITFDKGGTW EFLQAPAFTGYGEKINCELSQGCSLHLAQRLSQLLNLQLRRMPILSKESAPGLIIATGSVGKNLASKTNVYISSS AGARWREALPGPHYYTWGDHGGIITAIAQGMETNELKYSTNEGETWKTFIFSEKPVFVYGLLTEPGEKSTVFTIF GSNKENVHSWLILQVNATDALGVPCTENDYKLWSPSDERGNECLLGHKTVFKRRTPHATCFNGEDFDRPVVVSNC SCTREDYECDFGFKMSEDLSLEVCVPDPEFSGKSYSPPVPCPVGSTYRRTRGYRKISGDTCSGGDVEARLEGELV PCPLAEENEFILYAVRKSIYRYDLASGATEQLPLTGLRAAVALDFDYEHNCLYWSDLALDVIQRLCLNGSTGQEV IINSGLETVEALAFEPLSQLLYWVDAGFKKIEVANPDGDFRLTIVNSSVLDRPRALVLVPQEGVMFWTDWGDLKP GIYRSNMDGSAAYHLVSEDVKWPNGISVDDQWIYWTDAYLECIERITFSGQQRSVILDNLPHPYAIAVFKNEIYW DDWSQLSIFRASKYSGSQMEILANQLTGLMDMKIFYKGKNTGSNACVPRPCSLLCLPKANNSRSCRCPEDVSSSV LPSGDLMCDCPQGYQLKNNTCVKEENTCLRNQYRCSNGNCINSIWWCDFDNDCGDMSDERNCPTTICDLDTQFRC QESGTCIPLSYKCDLEDDCGDNSDESHCEMHQCRSDEYNCSSGMCIRSSWVCDGDNDCRDWSDEANCTAIYHTCE ASNFQCRNGHCIPQRWACDGDTDCQDGSDEDPVNCEKKCNGFRCPNGTCIPSSKHCDGLRDCSDGSDEQHCEPLC $\verb|Thfmdfvcknrqqclfhsmvcdgiiqcrdgsdedaafagcsqdpefhkvcdefgfqcqngvcisliwkcdgmddc|\\$ GDYSDEANCENPTEAPNCSRYFQFRCENGHCIPNRWKCDRENDCGDWSDEKDCGDSHILPFSTPGPSTCLPNYYR CSSGTCVMDTWVCDGYRDCADGSDEEACPLLANVTAASTPTQLGRCDRFEFECHQPKTCIPNWKRCDGHQDCQDG $\verb|RDEANCPTHSTLTCMSREFQCEDGEACIVLSERCDGFLDCSDESDEKACSDELTVYKVQNLQWTADFSGDVTLTW|\\$ MRPKKMPSASCVYNVYYRVVGESIWKTLETHSNKTNTVLKVLKPDTTYQVKVQVQCLSKAHNTNDFVTLRTPEGL PDAPRNLQLSLPREAEGVIVGHWAPPIHTHGLIREYIVEYSRSGSKMWASQRAASNFTEIKNLLVNTLYTVRVAA VTSRGIGNWSDSKSITTIKGKVIPPPDIHIDSYGENYLSFTLTMESDIKVNGYVVNLFWAFDTHKOERRTLNFRG SILSHKVGNLTAHTSYEISAWAKTDLGDSPLAFEHVMTRGVRPPAPSLKAKAINQTAVECTWTGPRNVVYGIFYA TSFLDLYRNPKSLTTSLHNKTVIVSKDEQYLFLVRVVVPYQGPSSDYVVVKMIPDSRLPPRHLHVVHTGKTSVVI KWESPYDSPDQDLLYAIAVKDLIRKTDRSYKVKSRNSTVEYTLNKLEPGG

834/2825 FIGURE 733

GGCACGAGGCTGCGCCGGAAGTCCCTAGCCAGGCCTGGCGGTAACCTTGGGGGCCTCACTGCAGCCGCCGCTGCT GTTGGAGTGGGCTTTGCGAGTCTGAACGTTGGCGGGGCTAGGCTCGTTAACTGCCGAGAGCCTCCGGGTTTGCGG $\tt TGGAGGACGCTGAGGCCGTGGGGGGCAGGCACCCGGGCGCGGGCCTCCCAGCCGACAT{\color{red} G}CTCTCTAGTGGCGGAC{\color{red} G}CGGAC{\color{red} G}CGGAC{{\color{red} G}CGGCGAC{{\color{red} G}CGGAC{{\color{red} G}CGGCAC{{\color{red} G}CGGAC{{\color{red} G}CG$ AGCCTTCGTCTCCCAGATTGCAGCTGCAGAACCTTGGCCTGAAAATGCTACATTATATCAGCAATTGAAAGGGGA GCAAATTTTACTTTCTGACAATGCAGCTTCTCTTGCAGTGCAGGCCTTTTTGCAAATGTGTAACTTGCCTATCAA AGTAGTTTGTAGGGCAAATGCAGAATATATGTCTCCATCTGGTAAAGTACCTTTTATTCATGTGGGAAATCAAGT AGTATCAGAACTTGGTCCAATAGTCCAATTTGTTAAAGCCAAGGGCCATTCTCTTAGTGATGGGCTGGAGGAAGT $\verb|CCAAAAAGCAGAAATGAAAGCTTACATGGAATTAGTCAACAATATGCTGTTGACTGCAGAGCTGTATCTTCAGTG|$ GTGTGATGAAGCTACAGTAGGGGAGATCACTCATGCTAGGTATGGATCTCCTTACCCTTGGCCTCTGAATCATAT TTTGGCCTATCAAAAACAGTGGGAAGTCAAACGTAAGATGAAAGCTATTGGATGGGGAAAGAAGACTCTGGACCA GGTCTTAGAGGATGTAGACCAGTGCTGTCAAGCTCTCTCAAAGACTGGGAACACCAACCGTATTTCTTCAATAA GCAGCCTACTGAACTTGACGCACTGGTATTTGGCCATCTATACACCATTCTTACCACACAATTGACAAATGATGA ACTTTCTGAGAAGGTGAAAAACTATAGCAACCTCCTTGCTTTCTGTAGGAGAATTGAACAGCACTATTTTGAAGA ${\tt TCGTGGTAAAGGCAGGCTGTCA{\tt TAG}GGTTATGTGTTAGTCTCAGGAGTCTTAACTTTTGAAATATGTTTTACTTG}$ ATCTTATGTATGTGCTTTATATTGTTATTTGTGTATACATTAAAATAATTCTGAATTATTTAATCTGATATGTTG

835/2825 FIGURE 734

 ${\tt MSLVAEAFVSQIAAAEPWPENATLYQQLKGEQILLSDNAASLAVQAFLQMCNLPIKVVCRANAEYMSPSGKVPFI$$ HVGNQVVSELGPIVQFVKAKGHSLSDGLEEVQKAEMKAYMELVNNMLLTAELYLQWCDEATVGEITHARYGSPYP$$ WPLNHILAYQKQWEVKRKMKAIGWGKKTLDQVLEDVDQCCQALSQRLGTQPYFFNKQPTELDALVFGHLYTILTTQLTNDELSEKVKNYSNLLAFCRRIEQHYFEDRGKGRLS$

836/2825 FIGURE 735A

GCGGCCGCGTCGACGCGGCGGCGGCGCGCGTCGGCTCGGGGTTCTCCGGGAGAGGGGGAGTGCGCGGCGGCCG CAGCTGCCACAAACCAGGTGAAGCTTTGTTCTAAGAATATTTGTTTCATCTAGTTTATGAGTCCAAATGATATAG ACTGTAAATGTCACAGCAGTGGTGAAAGACTGCTCGGTCATGAGCACCGACAGTAACTCACTGGCACGTGAATTT CTGACCGATGTCAACCGGCTTTGCAATGCAGTGGTCCAGAGGGTGGAGGCCAGGGAGGAAGAAGAGAGGAGAACA GATCAGGCATTGACATGTAGAGAAGAACTCCTGACTCTTCTTCTGTCTCTCCTTCCACTGGTATGGAAGATACCT GTCCAAGAAGAAAAGGCAACAGATTTTAACCTACCGCTCTCAGCAGATATAATCCTGACCAAAGAAAAGAACTCA CTCTCCACCTCAGATTCAGAAGCCAATTCAGATGAAAAAGGCATAGCAATGAATAAGCATAGAAGGCCCCATCTG CTGCATCATTTTTTAACATCGTTTCCTAAACAAGACCACCCCAAAGCTAAACTTGACCGCTTAGCAACCAAAGAA CAGACTCCTCCAGATGCTATGGCTTTGGAAAATTCCAGAGAGATTATTCCAAGACAGGGGTCAAACACTGACATT TTAAGTGAGCCAGCTGCCTTGTCTGTTATCAGTAACATGAACAATTCTCCATTTGACTTATGTCATGTTTTGTTA TCTTTATTAGAAAAGTTTGTAAGTTTGACGTTACCTTGAATCATAATTCTCCTTTAGCAGCCAGTGTAGTGCCC ACACTAACTGAATTCCTAGCAGGCTTTGGGGGACTGCTGCAGTCTGAGCGACAACTTGGAGAGTCGAGTAGTTTCT GCAGGTTGGACCGAAGAACCGGTGGCTTTGATTCAAAGGATGCTCTTTCGAACAGTGTTGCATCTTCTGTCAGTA GATGTTAGTACTGCAGAGATGATGCCAGAAAATCTTAGGAAAAATTTAACTGAATTGCTTAGAGCAGCTTTAAAA ATTAGAATATGCCTAGAAAAGCAGCCTGACCCTTTTGCACCAAGACAAAAGAAAACACTGCAGGAGGTTCAGGAA GATTTTGTGTTTTCAAAGTATCGTCATAGAGCCCTTCTTTTACCTGAGCTTTTTGGAAGGAGTTCTTCAGATTCTG ${\tt ATCTGTTGTCTTCAAAGTGCAGCTTCAAATCCCTTCTACTTCAGTCAAGCCATGGATTTGGTTCAAGAATTCATT}$ ${\tt CAGCATCATGGATTTAATTTATTTGAAACAGCAGTTCTTCAAATGGAATGGCTGGTTTTAAGAGATGGAGTTCCT}$ CCCGAGGCCTCAGAGCATTTGAAAGCCCTAATAAATAGTGTGATGAAAATAATAGTGAGCACTGTCAAAAAAGTGAAA TCAGAGCAACTTCATCATTCGATGTGTACAAGAAAAAGGCACAGACGATGTGAATATTCTCATTTTATGCATCAT CACCGAGATCTCTCAGGTCTTCTGGTTTCGGCTTTTAAAAACCAGGTTTCCAAAAACCCATTTGAAGAGACTGCA GATGGAGATGTTTATTATCCTGAGCGGTGCTGTTGCATTGCAGTGTGTGCCCATCAGTGCTTGCGCTTACTACAG CAGGCTTCCTTGAGCAGCACTTGTGTCCAGATCCTATCGGGTGTTCATAACATTGGAATATGCTGTTGTATGGAT CCCAAATCTGTAATCATTCCTTTGCTCCATGCTTTTAAATTGCCAGCACTGAAAAAATTTTCAGCAGCATATATTG AATATCCTTAACAAACTTATTTTGGATCAGTTAGGAGGAGCAGAGATATCACCAAAAATTAAAAAAGCAGCTTGT AATATTTGTACTGTTGACTCTGACCAACTAGCCCAATTAGAAGAGACACTGCAGGGAAACTTATGTGATGCTGAA CTCTCCTCAAGTTTATCCAGTCCTTCTTACAGATTTCAAGGGATCCTGCCCAGCAGTGGATCTGAAGATTTGTTG TGGAAATGGGATGCTTTAAAGGCTTATCAGAACTTTGTTTTTTGAAGAAGACAGATTACATAGTATACAGATTGCA AATCACATTTGCAATTTAATCCAGAAAGGCAATATAGTTGTTCAGTGGAAATTATATAATTACATATTTAATCCT AGCCATCATAACCAGTGCTTGCCTCAGGACGTGCTTCAGATTTATGTAAAAACTCTGCCTATCCTGCTTAAATCC AGGGTAATAAGAGATTTGTTTTTGAGTTGTAATGGAGTAAGTCAAATAATCGAATTAAATTGCTTAAATGGTATT CGAAGTCATTCTCTAAAAGCATTTGAAACTCTGATAATCAGCCTAGGGGAGCAACAGAAAGATGCCTCAGTTCCA GATATTGATGGGATAGACATTGAACAGAAGGAGTTGTCCTCTGTACATGTGGGTACTTCTTTTCATCATCAGCAA GCTTATTCAGATTCTCCTCAGAGTCTCAGCAAATTTTATGCTGGCCTCAAAGAAGCTTATCCAAAGAAGACGGAAG AGTCATATGCTGCCATGTATATCTCTCGAGAGCCTTGTCTTGCCTTCTCCTGAACATATGCACCAAGCAGCAGAC ATTTGGTCTATGTGTCGTTGGATCTACATGTTGAGTTCAGTGTTCCAGAAACAGTTTTATAGGCTTGGTGGTTTC CGAGTATGCCATAAGTTAATATTTATGATAATACAGAAACTGTTCAGAAGTCACAAAGAGGAGCAAGGAAAAAAG TTATTATCTTTGGCTATAAAAAGTGACCCCATACCATCAGAACTAGGTAGTCTAAAAAAAGAGTGCTGACAGTTTA GGTAAATTAGAGTTACAGCATATTTCTTCCATAAATGTGGAAGAAGTTTCAGCTACTGAAGCCGCTCCCGAGGAA GCAAAGCTATTTACAAGTCAAGAAAGTGAGACCTCACTTCAAAGTATACGACTTTTGGAAGCCCTTCTGGCCATT TGTCTTCATGGTGCCAGAACTAGTCAACAGAAGATGGAATTGGAGTTACCTAATCAGAACTTGTCTGTGGAAAGT

837/2825 FIGURE **735B**

ATATTATTTGAAATGAGGGACCATCTTTCCCAGTCAAAGGTGATTGAAACACAACTAGCAAAGCCGTTATTTGAT GCCCTGCTTCGAGTTGCCCTCGGGAATTATTCAGCAGATTTTGAACATAATGATGCTATGACTGAGAAGAGTCAT CAATCTGCAGAAGAATTGTCATCCCAGCCTGGTGATTTTTCAGAAGAAGCTGAGGATTCTCAGTGTTGTAGTTTT AAACTTTTAGTTGAAGAAGAAGGTTACGAAGCAGATAGTGAAAGCAATCCTGAAGATGGCGAAACCCAGGATGAT GGGGTAGACTTAAAGTCTGAAACAGAAGGTTTCAGTGCATCAAGCAGTCCAAATGACTTACTCGAAAACCTCACT ${\tt CAAGGGGAAATAATTTATCCTGAGATTTGTATGCTGGAATTAAATTTGCTTTCTGCTAGTAAAGCCAAACTTGAT}$ CAGGGAACTGTGAAAAATCTTTTAGGAGGGTTCTTGAGTATTTTAACACAGGATGATTCTGATTTTCAAGCATGC AGAATATTTCTGGAGAAATCTCCTTGTACAAAAATTCTTCTTCTGGGTATTCTGAAAATTATTGAAAGTGATACT ACTATGAGCCCTTCACAGTATCTAACCTTCCCTTTACTGCACGCTCCAAATTTAAGCAACGGTGTTTCATCACAA GCTGATAGAGAGAGTTTTCCCCATCGGCTGCTTTCATCTTGGCACATAGCCCCAGTCCACCTGCCGTTGCTGGGG CAAAACTGCTGGCCACCCTATCAGAAGGTTTCAGTGTTTTCCCTGTGGTTTAATGTGGAGTGTATCCATGAAGCT GGTACAGAGAGCGACAGACCAGAAGGTGCAGAGTACATAAATCCTGGTGAAAGACTCATAGAAGAAGGATGTATT CATATAATTTCACTGGGATCCAAAGCGTTGATGATCCAAGTGTGGGCTGATCCCCACAATGCCACTCTTATCTTT CGTGTGTGCATGGATTCAAATGATGACATGAAAGCTGTTTTACTAGCACAGGTTGAATCACAGGAGAATATTTTC CTCCCAAGCAAATGGCAACATTTAGTACTCACCTACTTACAGCAGCCCCCAAGGGAAAAGGAGGATTCATGGGAAA ATCTCCATATGGGTCTCTGGACAGAGGAAGCCTGATGTTACTTTGGATTTTATGCTTCCAAGAAAAACAAGTTTG TCATCTGATAGCAATAAAACATTTTGCATGATTGGCCATTGTTTATCATCCCAAGAAGAGTTTTTGCAGTTGGCT GCTTGTGGACCCAACCATACATCTGTAATGCCATGTAAGTATGGCAAGCCAGTCAATGACTACTCCAAATATATT ATTGAAAGTCTTTCAGTTGTTTATACAACTTACTGTCCTGCTCAGTATACCATCTATGAACCAGTGATTAGACTT AAAGGTCAAATGAAAACCCAACTCTCTCAAAGACCCTTCAGCTCAAAAGAAGTTCAGAGCATCTTATTAGAACCT CATCATCTAAAGAATCTCCAACCTACTGAATATAAAACTATTCAAGGCATTCTGCACGAAATTGGTGGAACTGGC ATATTTGTTTTTCTCTTTGCCAGGGTTGTTGAACTCAGTAGCTGTGAAGAAACTCAAGCATTAGCACTGCGAGTT ATACTCTCATTAATTAAATACAACCAACAAAGAGTACATGAATTAGAAAATTGTAATGGACTTTCTATGATTCAT CAGGTGTTGATCAAACAAAAATGCATTGTTGGGTTTTACATTTTGAAGACCCTTCTTGAAGGATGCTGTGGTGAA GATATTATTTATGAATGAGAATGGAGAGTTTAAGTTGGATGTAGACTCTAATGCTATAATCCAAGATGTTAAG GCTCTAGAAGTCCTCATCAGAGCAGATCACCACCAGCAGATGTTTAATATTAAGCAGTTATTGAAAGCTCAAGTG GTTCATCACTTTCTACTGACTTGTCAGGTTTTGCAGGAATACAAAGAGGGGCCAACTCACACCCATGCCCCGAGAG GTTTGTAGATCATTTGTGAAAATTATAGCAGAAGTCCTTGGATCTCCTCCAGATTTGGAATTATTGACAATTATC TTCAATTTCCTTTTAGCAGTTCACCCTCCTACTAATACTTACGTTTGTCACAATCCCACGAACTTCTACTTTTCT TTGCACATAGATGGCAAGATCTTTCAGGAGAAAGTGCGGTCAATCATGTACCTGAGGCATTCCAGCAGTGGAGGA AGGTCCCTTATGAGCCCTGGATTTATGGTAATAAGCCCATCTGGTTTTACTGCTTCACCATATGAAGGAGAGAAT TCCTCTAATATTATTCCACAACAGATGGCCGCCCATATGCTGCGTTCTAGAAGCCTACCAGCATTCCCTACTTCT TCACTACTAACGCAATCACAAAAACTGACTGGAAGTTTGGGTTGTAGTATCGACAGGTTACAAAATATTGCAGAT ACTTATGTTGCCACCCAATCAAAGAAACAAAATTCTTTGGGGAGTTCCGACACACTGAAAAAAGGCAAAGAGGAC GCATTCATCAGTAGCTGTGAGTCTGCAAAAACTGTTTGTGAAATGGAAGCTGTCCTCTCAGCCCAGGTCTCTGTC AGTGATGTCCCAAAGGGAGTGCTGGGATTTCCAGTGGTCAAAGCAGATCATAAACAGTTGGGAGCAGAACCCAGG TCAGAAGATGACAGTCCTGGGGATGAGTCCTGCCCACGCCGACCTGATTACCTAAAGGGATTGGCCTCCTTCCAG CGAAGCCACAGCACTATTGCAAGCCTTGGGCTAGCTTTTCCTTCACAGAACGGATCTGCAGCTGTTGGCCGTTGG CCAAGTCTTGTTGATAGAAACACTGATGATTGGGAAAACTTTGCCTATTCTCTTGGTTATGAGCCAAATTACAAC CGAACTGCAAGTGCTCACAGTGTAACTGAAGACTGTTTGGTACCTATATGCTGTGGATTATATGAACTCCTAAGT GGGGTTCTTCTTATCCTGCCTGATGTTTTGCTTGAAGATGTGATGGACAAGCTTATTCAAGCAGATACACTTTTG

838/2825 FIGURE **735**C

GTCCTCGTTAACCACCCATCACCAGCTATACAACAAGGTGTTATTAAACTATTAGATGCATATTTTGCTAGAGCA TCTAAGGAACAAAAGATAAATTTCTGAAGAATCGTGGATTTTCCTTGCTAGCCAACCAGTTGTATCTTCATCGA GGAACTCAAGAATTGTTAGAATGCTTCATCGAAATGTTCTTTGGTCGACATATTGGCCTTGATGAAGAATTTTGAT CTGGAAGATGTGAGAAACATGGGATTGTTTCAGAAGTGGTCTGTCATTCCTATTCTGGGACTAATAGAGACCTCT CTATATGACAACATACTCTTGCATAATGCTCTTTTACTTCTTCTCCAAATTTTAAATTCTTGTTCTAAGGTAGCA GATATGTTGCTGGATAATGGTCTACTCTATGTGTTATGTAATACAGTAGCAGCCCTGAATGGATTAGAAAAGAAC ATTCCCATGAGTGAATATAAATTGCTTGCTTGTGATATACAGCAACTTTTCATAGCAGTTACAATTCATGCTTGC AGTTCCTCAGGCTCACAATATTTTAGGGTTATTGAAGACCTTATTGTAATGCTTGGATATCTTCAAAATAGCAAA AACAAGAGGACACAAAATATGGCTGTTGCACTACAGCTTAGAGTTCTCCAGGCTGCTATGGAATTTATAAGGACC ACCGCAAATCATGACTCTGAAAACCTCACAGATTCACTCCAGTCACCTTCTGCTCCCCATCATGCAGTAGTTCAA AAGCGGAAAAGCATTGCTGGTCCTCGAAAATTTCCCCTTGCTCAAACTGAATCGCTTCTGATGAAAATGCGTTCA GTGGCAAATGATGAGCTTCATGTGATGATGCAACGGAGAATGAGCCAAGAAACCCTAGCCAAGCAACTGAAACG ATTATTGACATTTTGAGAACTCCAGAAAATGTAACTCAAAGCAAGACCTCAGTTTTCCAGACCGAAATTTCTGAG TTCAAAGTATCTATTGGTTCAAGTAAAGCCAGTGGTTCCAAGCAGCAATGGACTAAAATTCTGTGGTCTTGTAAG GAGACCTTCCGAATGCAGCTTGGGAGACTACTAGTGCATATTTTGTCGCCAGCCCACGCTGCACAAGAGAGAAAG CAAATTTTTGAAATAGTTCATGAACCAAATCATCAGGAAATACTACGAGACTGTCTCAGCCCCATCCCTACAACAT GGAGCCAAGTTAGTTTTGTATTTGTCAGAGTTGATACATAATCACCAAGGTGAATTGACTGAAGAAGAGCTAGGC GACCTTATTAAAATGATCAAAGAGGAACAAAAGAAATATGAAACTGAAGAAGGAGTGAATAAAGCTGCTTGGCAG AAAACAGTTAACAATAATCAACAAAGTCTCTTTCAGCGTCTGGATTCAAAATCAAAGGATATATCTAAAATAGCT GCAGATATCACCCAGGCAGTGTCTCTCTCCCAAGGAAATGAGAGAAAAAAGGTGATCCAGCATATTAGAGGAATG TATAAAGTAGATTTGAGTGCCAGCAGACATTGGCAGGAACTTATTCAGCAGCTGACACATGATAGAGCAGTATGG CAGAGATGTTATTTAACTATTCCAAATAAGTATCTCCTTAGGGATAGACAGAAATCAGAAGATGTTGTCAAACCA CCACTCTCTTACCTGTTTGAAGACAAAACTCATTCTTCTTCTTCTTCTACTGTCAAAGACAAAGCTGCAAGTGAA TCTATAAGAGTGAATCGAAGATGCATCAGTGTTGCACCATCTAGAGAGACAGCTGGTGAATTGTTACTAGGTAAA TGTGGAATGTATTTTGTGGAAGATAATGCTTCTGATACAGTTGAAAGTTCGAGCCTTCAGGGAGAGTTGGAACCA GCATCATTTTCCTGGACATATGAAGAAATTAAAGAAGTTCACAAGCGTTGGTGGCAATTGAGAGATAATGCTGTA GAAATCTTTCTAACAAATGGCAGAACACTCCTGTTGGCATTTGATAACACCAAGGTTCGTGATGATGTATACCAC AATATACTCACAAATAACCTCCCTAATCTTCTGGAATATGGTAACATCACCGCTCTGACAAATTTATGGTATACT TATCCTGTGTTCCCATTTATACTTGCTGACTACGTTAGTGAGACACTTGACCTCAATGATCTGTTGATATACAGA AATCTCTCTAAACCTATAGCTGTTCAGTATAAAGAAAAAGAAGATCGTTATGTGGACACATACAAGTACTTGGAG GAAGAGTACCGCAAAGGAGCCAGAGAAGATGACCCCATGCCTCCCGTGCAGCCCTATCACTATGGCTCCCACTAT TCCAATAGCGGCACTGTGCTTCACTTCCTGGTCAGGATGCCTCCTTTCACTAAAATGTTTTTAGCCTATCAAGAT CAAAGTTTTGACATTCCAGACAGAACTTTTCATTCTACAAATACAACTTGGCGACTCTCATCTTTTGAATCTATG ACTGATGTGAAAGAACTTATCCCAGAGTTTTTCTATCTTCCAGAGTTCCTAGTTAACCGTGAAGGTTTTGATTTT GGTGTGCGTCAGAATGGTGAACGGGTTAATCACGTCAACCTTCCCCCTTGGGCGCGTAATGATCCTCGTCTTTTT ATCCTCATCCATCGGCAGGCTCTAGAGTCTGACTACGTGTCGCAGAACATCTGTCAGTGGATTGACTTGGTGTTT GGGTATAAGCAAAAGGGGAAGGCTTCTGTTCAAGCGATCAATGTTTTTCATCCTGCTACATATTTTGGAATGGAT GTCTCTGCAGTTGAAGATCCAGTTCAGAGACGAGCGCTAGAAACCATGATAAAAACCTACGGGCAGACTCCCCGT CAGCTGTTCCACATGGCCCATGTGAGCAGACCTGGAGCCAAGCTCAATATTGAAGGAGAGCTTCCAGCTGCTGTG GGGTTGCTAGTGCAGTTTGCTTTCAGGGAGACCCGAGAACAGGTCAAAGAAATCACCTATCCGAGTCCTTTGTCA TGGATAAAAGGCTTGAAATGGGGGGAATACGTGGGTTCCCCCAGTGCTCCAGTACCTGTGGTCTGCTTCAGCCAG CCCCACGGAGAAAGATTTGGCTCTCCCAGGCTCTGCCCACCAGAGCAATCTGTGGTTTGTCACGGAATTTCTGT CTTGTGATGACATATAGCAAGGAACAAGGTGTGAGAAGCATGAACAGTACGGACATTCAGTGGTCAGCCATCCTG

839/2825 FIGURE **735D**

AGCTGGGGATATGCTGATAATATTTTAAGGTTGAAGAGTAAACAAAGTGAGCCTCCAGTAAACTTTATTCAAAGT TCACAACAGTACCAGGTGACTAGTTGTGCTTGGGTGCCTGACAGTTGCCAGCTGTTTACTGGAAGCAAATGCGGT GTCATCACAGCCTACACAAACAGATTTACAAGCAGCACGCCATCAGAAATAGAAATGGAGACTCAAATACATCTC TATGGTCACACAGAAGAGATAACCAGCTTATTTGTTTGCAAACCATACAGTATACTGATAAGTGTGAGCAGAGAC GGAACCTGCATCATATGGGATTTAAACAGGTTATGCTATGTACAAAGTCTGGCGGGACACAAAAGCCCTGTCACA CTCTGGACGGTGAACGGGGATCTCGTTGGACATGTCCACTGCAGGGAGATCATCTGTTCCGTGGCTTTCTCCAAC CAGCCTGAGGGAGTATCTATCAATGTAATCGCTGGGGGATTAGAAAATGGAATTGTAAGGTTATGGAGCACATGG GACTTAAAGCCTGTGAGAGAAATTACATTTCCCAAATCAAATAAGCCCATCATCAGCCTTACATTTTCTTGTGAT GGCCACCATTTGTACACAGCAAACAGTGATGGGACCGTGATTGCCTGGTGTCGGAAGGACCAGCAGCGCTTGAAA CAGCCAATGTTCTATTCCTTCCTTAGCAGCTATGCAGCCGGGTGAATGCGAATGAACTTCACGTTCTCCAAAGCA CTTTAACTCCAAACTAGATTTGTTGACTTCACCAGTTTTAGGAGGTTGAACCTAAAGAAATGGATGACTGGACAA ACCATCCAAATAATGATAAAGTCTATTCATCTGCACAAAATTCTGAAGAGTCACATGATCCTAAGAGGAAAGTTC TGTTCTATTTTAGTGATAATCTGGAAGATTGTGTCAATATGCACTAGCCAACAAGTTTTAAGCCTCGCATGGTAC ATTAAAATGATATTCTTAAAATTTTTTCCCACCAAGGTATTCCAAAGAAAATATTAAGGTCTCCCCTTTTTCTAT GATTCCAAAAGGACCAGTAGAATTTAAATTGGTTGGTTGATXGTTTATATAAAACACACTAAAATTATATTTTAA AGGACTTGATAATTGCTTTCTTTGAGAAGCAAATTATTCAGTAGGTGCCTCTGTACCAAATATTTTATGGAATAT CTAAATACTAAAATAAACTATGAATGAATCTCAAAATTAGGCAGTTTTTTGCCAGTTGCTTTCTTAGCTCAAAGGA GAACCAGAATTTTTTTGACAGCCACAAACAAGAATACAGGTATCTTGGATTTCAGACACATTCTGTTTCTTCATA AAAATTTTACTTAAAATCTGTAACGCTAGATATTGACTATCCTTAGTTGAGTCACTGAGGTTTAAACACAATGGT AAGTCTTAAAGTCTGCTATTTACAGAGCATTGAATCTGTACCAATTTGCAATAGAAAGCCTTCAGTATGCAAGAA GTTTGCATGGGTATTAAGAACACAGCCTAAATAAGGCATTTGATCTAATCTGCAGGAAGAATTTTCTTCCCCAAA ACAGAATTATAAAAGCTTACTTTAAACAGGAGGCAGAATAATTCTTTTAGGAAACCATTTCATTCTGTTTCTACT TTACTTGGGGTGCTGCTTTATTGGCTTTGAAAACACTGTCAGATAAGCTCAGTAATATGTTACCATGGGATAAAA ATATGTATCCCTGCCTAAGAATAACTTGTGCATTTGTTATGGAAATTTAATTCATATGGTGTTTACAGTACTACT TTTGTAACTTCCAGACTTTCTAAAACATTCTGCTTAAAAACCATATAAAATATAATTCCAAAGTCTCTGCTGTCA AGATAGATTCGAGAGAAAGCACGTGGCCATGTATGCTTTAACCTTAAACTGCATACACATGTAGTGATACCTAGG $\tt CTGCATTIAGATCACCGTGTGCTCAGGCCAGGTGTGAATCCTGAGGTCCATGGAGGTGCAGAGATGAGATTACTC$ CTATTCACGTTGAAGTGATTTGCTTTGTTAACAAAAATTGCAGCTATTGTCTAGCTTTCATTTTTTTACTGAGA ACTITAAATTAGTCCCCTATTAGAATAGGGTTGCTACTCATCTTTTTTTAAAAAACCGAATTTCATCATTTATCTA TGGGGGGATTATTTATTCAGCATGAAACCTAATATGTATATGTTTGAAATACTTCATAATGTGCATGTTGTAGCA AAAAAAAAAAAAAAAAAAAAAAA

840/2825 FIGURE 736

MSTDSNSLAREFLTDVNRLCNAVVQRVEAREEEEEETHMATLGQYLVHGRGFLLLTKLNSIIDQALTCREELLTL LLSLLPLVWKIPVOEEKATDFNLPLSADIILTKEKNSSSQRSTOEKLHLEGSALSSQVSAKVNVFRKSRRORKIT HRYSVRDARKTQLSTSDSEANSDEKGIAMNKHRRPHLLHHFLTSFPKQDHPKAKLDRLATKEQTPPDAMALENSR EIIPROGSNTDILSEPAALSVISNMNNSPFDLCHVLLSLLEKVCKFDVTLNHNSPLAASVVPTLTEFLAGFGDCC SLSDNLESRVVSAGWTEEPVALIQRMLFRTVLHLLSVDVSTAEMMPENLRKNLTELLRAALKIRICLEKQPDPFA PRQKKTLQEVQEDFVFSKYRHRALLLPELLEGVLQILICCLQSAASNPFYFSQAMDLVQEFIQHHGFNLFETAVL OMEWLVLRDGVPPEASEHLKALINSVMKIMSTVKKVKSEOLHHSMCTRKRHRRCEYSHFMHHHRDLSGLLVSAFK NQVSKNPFEETADGDVYYPERCCCIAVCAHQCLRLLQQASLSSTCVQILSGVHNIGICCCMDPKSVIIPLLHAFK LPALKNFQQHILNILNKLILDQLGGAEISPKIKKAACNICTVDSDQLAQLEETLQGNLCDAELSSSLSSPSYRFQ GILPSSGSEDLLWKWDALKAYONFVFEEDRLHSIQIANHICNLIOKGNIVVOWKLYNYIFNPVLQRGVELAHHCO HLSVTSAQSHVCSHHNQCLPQDVLQIYVKTLPILLKSRVIRDLFLSCNGVSQIIELNCLNGIRSHSLKAFETLII SLGEQQKDASVPDIDGIDIEQKELSSVHVGTSFHHQQAYSDSPQSLSKFYAGLKEAYPKRRKTVNQDVHINTINL FLCVAFLCVSKEAESDRESANDSEDTSGYDSTASEPLSHMLPCISLESLVLPSPEHMHQAADIWSMCRWIYMLSS VFQKQFYRLGGFRVCHKLIFMIIQKLFRSHKEEQGKKEGDTSVNENQDLNRISQPKRTMKEDLLSLAIKSDPIPS ELGSLKKSADSLGKLELQHISSINVEEVSATEAAPEEAKLFTSQESETSLQSIRLLEALLAICLHGARTSQQKME LELPNONLSVESILFEMRDHLSOSKVIETOLAKPLFDALLRVALGNYSADFEHNDAMTEKSHOSAEELSSOPGDF SEEAEDSQCCSFKLLVEEEGYEADSESNPEDGETQDDGVDLKSETEGFSASSSPNDLLENLTQGEIIYPEICMLE LNLLSASKAKLDVLAHVFESFLKIIRQKEKNVFLLMQQGTVKNLLGGFLSILTQDDSDFQACQRVLVDLLVSLMS SRTCSEELTLLLRIFLEKSPCTKILLLGILKIIESDTTMSPSOYLTFPLLHAPNLSNGVSSOKYPGILNSKAMGL LRRARVSRSKKEADRESFPHRLLSSWHIAPVHLPLLGQNCWPHLSEGFSVSLWFNVECIHEAESTTEKGKKIKKR NKSLILPDSSFDGTESDRPEGAEYINPGERLIEEGCIHIISLGSKALMIQVWADPHNATLIFRVCMDSNDDMKAV LLAQVESQENIFLPSKWQHLVLTYLQQPQGKRRIHGKISIWVSGQRKPDVTLDFMLPRKTSLSSDSNKTFCMIGH CLSSQEEFLQLAGKWDLGNLLLFNGAKVGSQEAFYLYACGPNHTSVMPCKYGKPVNDYSKYINKEILRCEQIREL FMTKKDVDIGLLIESLSVVYTTYCPAQYTIYEPVIRLKGQMKTQLSQRPFSSKEVQSILLEPHHLKNLQPTEYKT IQGILHEIGGTGIFVFLFARVVELSSCEETQALALRVILSLIKYNQQRVHELENCNGLSMIHQVLIKQKCIVGFY ILKTLLEGCCGEDIIYMNENGEFKLDVDSNAIIQDVKLLEELLLDWKIWSKAEQGVWETLLAALEVLIRADHHQQ MFNIKQLLKAQVVHHFLLTCQVLQEYKEGQLTPMPREVCRSFVKIIAEVL

841/2825 FIGURE **737A**

CTGAGCCGCATCTGCAATAGCACACTTGCCCGGCCACCTGCTGCCGTGAGCCCTTTGCTGCTGAAGCCCCTGGGGT ATCTACAACTACTCAGCCTTGCTGCTGAGCGAGGACAAGGACACCTTGTACATAGGTGCCCGGGAGGCGGTCTTC GCTGTGAACGCACTCAACATCTCCGAGAAGCAGCATGAGGTGTATTGGAAGGTCTCAGAAGACAAAAAAGCAAAA TGTGCAGAAAAGGGGAAATCAAAACAGACAGAGTGCCTCAACTACATCCGGGTGCTGCAGCCACTCAGCGCCACT TCCCTTTACGTGTGTGGGACCAACGCATTCCAGCCGGCCTGTGACCACCTGAACTTAACATCCTTTAAGTTTCTG GGGAAAAATGAAGATGGCAAAGGAAGATGTCCCTTTGACCCAGCACACACCTACACATCCGTCATGGTTGATGGA GAACTTTATTCGGGGACGTCGTATAATTTTTTGGGAAGTGAACCCATCATCTCCCGAAATTCTTCCCACAGTCCT CTGAGGACAGAATATGCAATCCCTTGGCTGAACGAGCCTAGTTTCGTGTTTGCTGACGTGATCCGAAAAAGCCCA GACAGCCCGACGGCGAGGATGACAGGGTCTACTTCTTCTTCACGGAGGTGTCTGTGGAGTATGAGTTTGTGTTC ACCTCCTTCCTGAAAGCCCGACTCATCTGCTCCCGGCCAGACAGCGGCTTGGTCTTCAATGTGCTGCGGGATGTC TTCGTGCTCAGGTCCCCGGGCCTGAAGGTGCCTGTGTTCTATGCACTCTTCACCCCACAGCTGAACAACGTGGGG CTGTCGGCAGTGTGCGCCTACAACCTGTCCACAGCCGAGGAGGTCTTCTCCCACGGGAAGTACATGCAGAGCACC ACAGTGGAGCAGTCCCACACCAAGTGGGTGCGCTATAATGGCCCGGTACCCAAGCCGCGGCCTGGAGCGTGCATC GACAGCGAGGCACGGCCCCAACTACACCAGCTCCTTGAATTTGCCAGACAAGACGCTGCAGTTCGTTAAAGAC CAGATCGTGGTGGACCGGACCCAGGCCCTGGATGGGACTGTCTATGATGTCATGTTTTGTCAGCACAGACCGGGGA GCTCTGCACAAAGCCATCAGCCTCGAGCACGCTGTTCACATCATCGAGGAGACCCAGCTCTTCCAGGACTTTGAG $\tt CCAGTCCAGACCCTGCTGTCTTCAAAGAAGGGCAACAGGTTTGTCTATGCTGGCTCTAACTCGGGCGTGGTC$ CAGGCCCGCTGGCCTTCTGTGGGAAGCACGGCACCTGCGAGGACTGTGTGCTGGCGCGGGACCCCTACTGCGCC TGGAGCCCGCCCACAGCGACCTGCGTGGCTCTGCACCAGACCGAGAGCCCCAGCAGGGGTTTGATTCAGGAGATG AGCGGCGATGCTTCTGTGCCCGGATAAAAGTAAAGGAAGTTACCGGCAGCATTTTTTCAAGCACGGTGGCACA GCGGAACTGAAATGCTCCCAAAAATCCAACCTGGCCCGGGTCTTTTGGAAGTTCCAGAATGGCGTGTTGAAGGCC GAGAGCCCCAAGTACGGTCTTATGGGCAGAAAAAACTTGCTCATCTTCAACTTGTCAGAAGGAGACAGTGGGGTG TACCAGTGCCTGTCAGAGGAGAGGGTTAAGAACAAAACGGTCTTCCAAGTGGTCGCCAAGCACGTCCTGGAAGTG GTGTTGGTGGCATCCACCCAAGGGTCTTCTCCCCCAACCCCAGCCGTGCAGGCCACCTCCTCCGGGGCCATCACC CTTCCTCCCAAGCCTGCGCCCACCGGCACATCCTGCGAACCAAAGATCGTCATCAACACGGTCCCCCAGCTCCAC TCGGAGAAAACCATGTATCTTAAGTCCAGCGACAACCGCCTCCTCATGTCCTCTTCCTCTTCTTTGTTCTC TTCCTCTGCCTCTTTTTCTACAACTGCTATAAGGGATACCTGCCCAGACAGTGCTTGAAATTCCGCTCGGCCCTA CTAATTGGGAAGAAGCCCAAGTCAGATTTCTGTGACCGTGAGCAGAGCCTGAAGGAGACGTTAGTAGAGCCA ATCACCAGCAAAGTCCCCACGGATAGGGAGGACTCACAGAGGATCGACGACCTTTCTGCCAGGGACAAGCCCTTT GACGTCAAGTGTGAGCTGAAGTTCGCTGACTCAGACGCAGATGGAGAC**TGA**GGCCGGCTGTGCATCCCCGCTGGT GCCTCGGCTGCGACGTGTCCAGGCGTGGAGAGTTTTGTGTTTCTCCTGTTCAGTATCCGAGTCTCGTGCAGTGCT GCGTAGGTTAGCCCGCATCGTGCAGACAACCTCAGTCCTCTTGTCTATTTTCTCTTGGGTTGAGCCTGTGACTTG GAAGAAGGCCCACGGGCGCACGGAGTTCCTGAGCCCTTTCTGTAGTGGGGGGAAAGGTGGCTGGACCTCTGTTGG CTGAGAAGAGCATCCCTTCAGCTTCCCCTCCCCGTAGCAGCCACTAAAAGATTATTTAATTCCAGATTGGAAATG ACATTTTAGTTTATCAGATTGGTAACTTATCGCCTGTTGTCCAGATTGGCACGAACCTTTTCTTCCACTTAATTA TATTTAGAAGAAGATGTATATCTTCCAGATTTTGTTATATATTTTGGCATAAAATACGGCTTACGTTGCTTAAGAT CCCACTCACCTTTTTCTTTTTTTTTTTTTTTTAACTTTATTCCTTGAGGGAAGCATTGTTTTTTGGAGAGAT TTTCTTTCTGTACTTCGTTTTACTTTTTTTTTTTAACTTTTACTCTCCGAAGAAGAGAGGACCTTCCCACATC CACGAGGTGGGTTTTGAGCAAGGGAAGGTAGCCTGGATGAGCTGAGTGGAGCCAGGCTGGCCCAGAGCTGAGATG

842/2825 FIGURE 737B

843/2825 FIGURE 738

MRMCTPIRGLLMALAVMFGTAMAFAPIPRITWEHREVHLVQFHEPDIYNYSALLLSEDKDTLYIGAREAVFAVNA LNISEKQHEVYWKVSEDKKAKCAEKGKSKQTECLNYIRVLQPLSATSLYVCGTNAFQPACDHLNLTSFKFLGKNE DGKGRCPFDPAHSYTSVMVDGELYSGTSYNFLGSEPIISRNSSHSPLRTEYAIPWLNEPSFVFADVIRKSPDSPD GEDDRVYFFFTEVSVEYEFVFRVLIPRIARVCKGDQGGLRTLQKKWTSFLKARLICSRPDSGLVFNVLRDVFVLR SPGLKVPVFYALFTPQLNNVGLSAVCAYNLSTAEEVFSHGKYMQSTTVEQSHTKWVRYNGPVPKPRPGACIDSEA RAANYTSSLNLPDKTLQFVKDHPLMDDSVTPIDNRPRLIKKDVNYTQIVVDRTQALDGTVYDVMFVSTDRGALHK AISLEHAVHIIEETQLFQDFEPVQTLLLSSKKGNRFVYAGSNSGVVQAPLAFCGKHGTCEDCVLARDPYCAWSPP TATCVALHQTESPSRGLIQEMSGDASVCPDKSKGSYRQHFFKHGGTAELKCSQKSNLARVFWKFQNGVLKAESPK YGLMGRKNLLIFNLSEGDSGVYQCLSEERVKNKTVFQVVAKHVLEVKVVPKPVVAPTLSVVQTEGSRIATKVLVA STQGSSPPTPAVQATSSGAITLPPKPAPTGTSCEPKIVINTVPQLHSEKTMYLKSSDNRLLMSLFLFFFVLFLCL FFYNCYKGYLPRQCLKFRSALLIGKKKPKSDFCDREQSLKETLVEPGSFSQQNGEHPKPALDTGYETEQDTITSK VPTDREDSQRIDDLSARDKPFDVKCELKFADSDADGD

844/2825 FIGURE 739

845/2825 FIGURE 740

 ${\tt MTCKMSQLERNIETIINTFHQYSVKLGHPDTLNQGEFKELVRKDLQNFLKKENKNEKVIEHIMEDLDTNADKQLS} \\ {\tt FEEFIMLMARLTWASHEKMHEGDEGPGHHHKPGLGEGTP}$

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846/2825 FIGURE 741A

 $\tt CACGTTGGGTGACATAATGGGGTTTTTTTAATTATAGATTCACACTGCATTTATTCATCACCCCTGTCCTCAT$ CCATAACTCAAATTTACTACCAGCAACACAAAATACAAAGATGTGTCCAGTTTCACTACAGCTCTTCGCGTTTAC TTGATTGGAAGGCTGGTTGCTGGGACTGTAGCGTTTGCAGGAAGTCACTTAACTGTTTGGGAGCTGGAAAACCGA AGCTGAAGTTCTCTTTTGCCATAGGAACGAGCGCAACTGACTAGGAAAGATGTGTCCCAAAGCTCCGCAAGCTGG AACGTGAGCCAGGAGGCCCGGACCGGCCACGGGACCGCGAGGCACTCCGAAAGTGTGCGGCTGCCCCTTCCCTGC CTCCCAGCTGTTACCCTTTTAAATGTCAGTGTTCGAGGCTGTAGGGGTAGCACGAGGCAGCGAAACGGAACAGTC GGATTGGCCGCACGCCTCAGTTCTAGACGCACCTCTCCACCGAAGCCGTTCTGACTGGCAGGGGGAGAAAGTAAA CAGAGTTGAATCACCCTCCCCACTGGCCAATTGGAGGGGGTTTGGTTTGTGACGTGATGGGATTCTGCGAAATTG AAAGTGTCTCTTAGACCTGGCGCGCGGGGCCCTCGCCACCCGCGTCGGGGTGATCGGGTGAATGTCCTGGGG CTTTGGCTCGACGCGAGGCCGAGGGCGTGCACCTCTCTTGCAGTTTCCTCTCCCAGCGCCTCGGGGGCGTT GTCGGGGGATCGCGGCCGGGGGCAGGGGCGGCGGCGGCGGCGGCGGGGGCCGGGGCCGTGGA TGCTGCGTGCGGAGGCGCTGCCGGTTACGTAAAGATGAGGGGCTGAGGTCGCCTCGGCGCTCCTGCGAGTCGGAA GAGCCGAACCTCCGCAGCCACCGCCAAGTTTGTCCGCGCCGCCTGGGCTGCCGTCGCCCGCACCATGTCCGCGGC CGCTCCGGACGCCGAGCGCTGCGACTACCTGAGCGCGAGGTGACCAAGGAGCACGGTGACCCGGGGGACACCTG GAAGGATTACTGCACACTGGTCACCATCGCCAAGAGCTTGTTGGACCTGAACAAGTACCGACCCATCCAGACCCC CTCCGTGTGCAGCGACAGTCTGGAAAGTCCAGATGAGGATATGGGATCCGACAGCGACGTGACCACCGAATCTGG TGGAGTGGCTGCGAAGGGGAAACACGCCTCCGAAAAGAGGCACAAGTGCCCCTACAGTGGCTGTGGGAAAGTCTA TGGAAAATCCTCCCATCTCAAAGCCCATTACAGAGTGCATACAGGTGAACGGCCCTTCCCCTGCACGTGGCCAGA CTGCCTTAAAAAGTTCTCCCGCTCAGACGAGCTGACCCGCCACTACCGGACCCACACTGGGGAAAAGCAGTTCCG CTGTCCGCTGTGTGAGAAGCGCTTCATGAGGAGTGACCACCTCACAAAGCACGCCCGGCGGCACACCGAGTTCCA CCCCAGCATGATCAAGCGATCGAAAAAGGCGCTGGCCAACGCTTTG**TGA**GGTGCTGCCCGTGGAAGCCAGGGAGG GATGGACCCCGAAAGGACAAAAGTACTCCCAGGAAACAGACGCGTGAAAACTGAGCCCCAGAAGAGGCACACTTG ACGGCACAGGAAGTCACTGCTCTTTGGTCAATATTCTGATTTTCCTCTCCCTGCATTGTTTTTAAAAAGCACATT GTAGCCTAAGATCAAAGTCAACACTCGGTCCCCTTGAAGAGGCAACTCTCTGAACCCGTCTCTGACTGTTGG GGGTAGGGAGTGGGTGGGGGGGGGGGGTAAGGCCAAGACTGGGTAGATTTTAAAGATTCAACACTGGTGTACAT TCTGTCATCTAAAGACTTGAAAACAAAAAACAGTTACTTATAGTCAATGGGTAAGCAGAGTCTGAATTTATACTA ATCAAGACAAACCTTTGAAAGGTTACACTAAGTACAGAACTTTTAAACCTTGCTTTGTATGAGTTGTACTTTTTG AACATAAGCTGCACTTTTATTTTCTAATGCAGAGGATGAATAAGTTAAATACATGCTTTGAGGATAGAAGCAGAT GTTCTGTTTGGCACCACGTTATAATCTGCTTATTTTACAATATACACGTTTCCCTAAGAAATCATGCGCAGAGAT AACAGAATTTTAACTCTATTAACTTTTCCAAATTTTCCTATGCTTTTAGTTAACATCATTATTGTATCCTAATGC CACTAGGGGAGAGGCTTTTGACTCTGTTGGGTTTTATTTGAATGTGTGCATAACAGTAATGAGATCTGGAAACA CCTATTTTTTGGGGAAAAAGGTTTGTTGGTCTCCTTCCTGTGTTCCTACAAAACTCCCACTCTCAGGTGCAAGAG TTATGTAGAAGGAAAGGGAGCTGAAATAGGAACAGAAAAATCAACCCCTATAACTAGTGAACACCAAGGGAAAAT ACCACATGATTCAGAGGAGACTCTGCAAAATCGTCCCTTGTGGAGAATGCAGGCAACATGGAATACTACGAAT GAAATCACATCACTGTATCTTTTACATCAATAGCCTCACCACTAATATATCTTGTATCTAGGTGTCTATAATGGC TGAAACCACTACATCCATCTATGCCATTTACCTGAAAACTTAACTGTGGCCTTTATGAGGCCAGAAAAGTGAACT

847/2825 FIGURE 741B

 ${\tt TCAGAAGTAACGCTTTCTGATGATTTTATCTGATATTTAAAACAGGGAGCTATGGTGCACTCTAGTTTATACTTG}$ CGCTCTGAAATGTGTAAACATAGGGTGCCTACCTATTTCACCTGACCCATACTCGTTTCTGATTCAGAATCAGTG TGGGCTCCTGCAGTGGGCGCGGGTCACGGCTGACTCCAACTTCCAATACAACAGCCATCACTAGCACAGTGTTTT TTTGTTTAACCAACGTAGTGTTATTAGTAGTTCTATAAAGAGAACTGCTTTTAACATTAGGGACTGGGAGCAGTC CATGGGATAAAAAGGAAAGTGTTTTCTCACGAGAAAACATGTCAGGAAAAATAAAGAACACTTTCTACCTCTGTT TCAGATTTTTGAAACACTTATTTTAAACCAAATTTTAATTTCTGTGTCCAAAATAAGTTTTAAGGACATCTGTTC TTCCATACGAAATAGGTTAGGCTGCCTATTTCTCACTGAGCTCATGGAATGGTTCTGCTTATGATACTCTGCACG $\verb|CTGCCTTTTAGTGAGTGAGGAGTTTGGGGTTGCCTAGCACTTGCTAACTTGTAAAAAGTCATCTTTCCCTCACAG| \\$ AAAGAAACGAAAGAAAGCAAAGTCAGTGAAAGACAATCTTTATAGTTTCAGGAGTAAATCTAAATGTGGC TGTTACTTGCCTTTTCAAGTGTGACAAACTCACATTTGATATTCTCTTATATGTTATAGTAATGTAACGTATAAA CTCAAGCCTTTTTATTCTTTGTGATTAAATCCTGTTTTAAAATGTCACAAAACAGGAACCAGCATTCTAATTAGA TTTACTATATCAAGATATGGTTCAAATAGGACTACTAGAGTTCATTGAACACTAAAACTATGAAACAATTACTTT TTATATTAAAAAGACCATGGATTTAACTTATGAAAATCCAAATGCAGGATAGTAATTTTTGTTTACTTTTTTAAC TTGTCATATTCTGGAAAATTTAATAGTTTTAGAGTTAAGATATCTCCTCTCTTTTGGTTAGGGAAGAAGAAAGCCC TTCACCATTGTGGAATGATGCCCTGGCTTTAAGGTTTAGCTCCACATCATGCTTCTCTT

848/2825 FIGURE 742

MSAAAYMDFVAAQCLVSISNRAAVPEHGVAPDAERLRLPEREVTKEHGDPGDTWKDYCTLVTIAKSLLDLNKYRP IQTPSVCSDSLESPDEDMGSDSDVTTESGSSPSHSPEERQDPGSAPSPLSLLHPGVAAKGKHASEKRHKCPYSGC GKVYGKSSHLKAHYRVHTGERPFPCTWPDCLKKFSRSDELTRHYRTHTGEKQFRCPLCEKRFMRSDHLTKHARRH TEFHPSMIKRSKKALANAL

849/2825 FIGURE **743**

GGCACGAGGGATCCCGGATGTGGATCAAGTTGGTGGGAAGCGTGCGGTGCCGCAGCA**ATC**GCGGCGCTCACAATT GCCACGGGTACTGGCAATTGGTTTTCGGCTTTGGCGCTCGGGGTGACTCTTCTCAAATGCCTTCTCATCCCCACA GCCAAATATTTTGATCAAGAAATGCTGAATGTCCATAATTTGAATTACTCCAGCTCAAGGACCTTACTTTTCCAG AGATTTTCCGTCATCTTTATGGATGTACTCTTTGTGTATGCTGTCCGTGAGTGCTGTAAATGCATTGATGGAAAA AAAGTGGGTAAAGAACTTACAGAAAAGCCAAAATTTATTCTGTCGGTATTACTTCTGTGGAACTTCGGGTTATTA ATTGTGGACCATATTCATTTTCAGTACAATGGCTTTTTATTTGGATTAATGCTACTCTCCATTGCACGATTATTT CAGAAAAGGCATATGGAAGGAGCATTTCTCTTTGCTGTTCTCCTACATTTCAAGCATATCTACCTCTATGTAGCA CCAGCTTATGGTGTATATCTGCTGCGATCCTACTGTTTCACTGCAAATAAACCAGATGGGTCTATTCGATGGAAG AGTTTCAGCTTTGTTCGTGTTATTTCCCTGGGACTGGTTGTTTTCTTAGTTTCTGCTCTTTCATTGGGTCCTTTC CTGGCCTTGAATCAGCTGCCTCAAGTCTTTTCCCGACTCTTTCCATTCAAGAGGGGCCCTCTGTCATGCATATTGG GTGACTCCCTTGGCAACCCTCATCTGCACACTGATTGCCATATTGCCCTCTATTTTCTGTCTTTGGTTTAAACCC CAAGGGCCCAGAGGCTTTCTCCGATGTCTAACTCTTTGTGCCTTGAGCTCCTTTATGTTTGGGTGGCATGTTCAT GAAAAAGCCATACTTCTAGCAATTCTCCCAATGAGCCTTTTGTCTGTGGGAAAAGCAGGAGACGCTTCGATTTTT CTGATTCTGACCACAACAGGACATTATTCCCTCTTTCCTCTGCTCTTCACTGCACCAGAACTTCCCATTAAAATC TTACTCATGTTACTATTCACCATATATAGTATTTCGTCACTGAAGACTTTATTCAGAAAAGAAAAACCTCTTTTT AATTGGATGGAAACTTTCTACCTGGCTTGGCCTGGGGCCTCTGGAAGTCTGCTGTGAATTTGTATTCCCTTTCACC TCCTGGAAGGTGAAGTACCCCTTCATCCCTTTGTTACTAACCTCAGTGTATTGTGCAGTAGGCGTCACATATGCT TGGTTCAAACTGTATGTTTCAGTATTGATTGACTCTGCTATTGGCAAGACAAAGAAACAA**TGA**ATAAAGGAACTG CTTAGAAAAAAAAAAAAAAAAAA

850/2825 FIGURE 744

MAALTIATGTGNWFSALALGVTLLKCLLIPTYHSTDFEVHRNWLAITHSLPISQWYYEATSEWTLDYPPFFAWFE
YILSHVAKYFDQEMLNVHNLNYSSSRTLLFQRFSVIFMDVLFVYAVRECCKCIDGKKVGKELTEKPKFILSVLLL
WNFGLLIVDHIHFQYNGFLFGLMLLSIARLFQKRHMEGAFLFAVLLHFKHIYLYVAPAYGVYLLRSYCFTANKPD
GSIRWKSFSFVRVISLGLVVFLVSALSLGPFLALNQLPQVFSRLFPFKRGLCHAYWAPNFWALYNALDKVLSVIG
LKLKFLDPNNIPKASMTSGLVQQFQHTVLPSVTPLATLICTLIAILPSIFCLWFKPQGPRGFLRCLTLCALSSFM
FGWHVHEKAILLAILPMSLLSVGKAGDASIFLILTTTGHYSLFPLLFTAPELPIKILLMLLFTIYSISSLKTLFR
KEKPLFNWMETFYLLGLGPLEVCCEFVFPFTSWKVKYPFIPLLLTSVYCAVGVTYAWFKLYVSVLIDSAIGKTKK

851/2825 FIGURE **745**

 $\verb|CCCCTCTTCCTCCTCAAGGGAAAGCTGCCCACTTCTAGCTGCCCTGCCATCCCCTTTAAAGGGCGACTTGCT|\\$ CAGCGCCAAACCGCGGCTCCAGCCCTCTCCAGCCTCCGGCTCAGCCGGCTCATCAGTCGGTCCGCGCCTTGCAGC TCCTCCAGAGGGACGCGCCCCGAG**ATG**GAGAGCCAAAGCCCTGCTGGTGACTCTGGCCGTGTGGCTCCAGAGT $\tt CTGACCGCCTCCCGCGGAGGGGTGGCCGCCGACCAAAGAAGAAGATTTTATCGACATCGAAAGTAAATTTGCC$ $\verb|CTAAGGACCCCTGAAGACACAGCTGAGGACACTTGCCACCTCATTCCCGGAGTAGCAGAGTCCGTGGCTACCTGT|\\$ CATTTCAATCACAGCAGCAAAACCTTCATGGTGATCCATGGCTGGACGGTAACAGGAATGTATGAGAGTTGGGTG CCAAAACTTGTGGCCGCCCTGTACAAGAGAGAACCAGACTCCAATGTCATTGTGGTGGACTGGCTGTCACGGGCT CAGGAGCATTACCCAGTGTCCGCGGGCTACACCAAACTGGTGGGACAGGATGTGGCCCGGTTTATCAACTGGATG GAGGAGGAGTTTAACTACCCTCTGGACAATGTCCATCTCTTGGGATACAGCCTTGGAGCCCATGCTGCTGGCATT GCAGGAAGTCTGACCAATAAGAAAGTCAACAGAATTACTGGCCTCGATCCAGCTGGACCTAACTTTGAGTATGCA GAAGCCCCGAGTCGTCTTTCTCCTGATGATGCAGATTTTGTAGACGTCTTACACACATTCACCAGAGGGTCCCCT GGTCGAAGCATTGGAATCCAGAAACCAGTTGGGCATGTTGACATTTACCCGAATGGAGGTACTTTTCAGCCAGGA AATAAAGTCAGAGCCAAAAGAAGCAGCAAAATGTACCTGAAGACTCGTTCTCAGATGCCCTACAAAGTCTTCCAT TACCAAGTAAAGATTCATTTTTCTGGGACTGAGAGTGAAACCCATACCAATCAGGCCTTTGAGATTTCTCTGTAT ATTTACACAGAGGTAGATATTGGAGAACTACTCATGTTGAAGCTCAAATGGAAGAGTGATTCATACTTTAGCTGG TCAGACTGGTGGAGCAGTCCCGGCTTCGCCATTCAGAAGATCAGAGTAAAAGCAGGAGAGACTCAGAAAAAGGTG ATCTTCTGTTCTAGGGAGAAGTGTCTCATTTGCAGAAAGGAAAGGCACCTGCGGTATTTGTGAAATGCCATGAC AAGTCTCTGAATAAGAAGTCAGGC**TGA**AACTGGGCGAATCTACAGAACAAAGAACGGCATGTGAATTCTGTGAAG AATGAAGTGGAGGAAGTAACTTTTACAAAACATACCCAGTGTTTTGGGGTGTTTCAAAAGTGGATTTTCCTGAATA TTAATCCCAGCCTTACCCTTGTTAGTTATTTTAGGAGACAGTCTCAAGCACTAAAAAGTGGCTAATTCAATTTAT GGGGTATAGTGGCCAAATAGCACATCCTCCAACGTTAAAAGACAGTGGATCATGAAAAGTGCTGTTTTGTCCTTT GAGAAAGAAATAATTGTTTGAGCGCAGAGTAAAATAAGGCTCCTTCATGTGGCGTATTGGGCCATAGCCTATAAT TGGTTAGAACCTCCTATTTTAATTGGAATTCTGGATCTTTCGGACTGAGGCCTTCTCAAACTTTACTCTAAGTCT TGATTCCCTCTTGCTATTGGAATGTGGTCCAGACGTCAACCAGGAACATGTAACTTGGAGAGGGACGAAGAAAGG GTCTGATAAACACAGAGGTTTTAAACAGTCCCTACCATTGGCCTGCATCATGACAAAGTTACAAATTCAAGGAGA TATAAAATCTAGATCAATTAATTCTTAATAGGCTTTATCGTTTATTGCTTAATCCCTCTCTCCCCCTTCTTTTTT GTCTCAAGATTATATATAATAATGTTCTCTGGGTAGGTGTTGAAAATGAGCCTGTAATCCTCAGCTGACACATA ATTTGAATGGTGCAGAAAAAAAAAAGATACCGTAATTTTATTATTATTAGATTCTCCAAATGATTTTCATCAATTTTAA AATCATTCAATATCTGACAGTTACTCTTCAGTTTTAGGCTTACCTTGGTCATGCTTCAGTTGTACTTCCAGTGCG $\verb|TCTCTTTGTTCCTGGCTTTGACATGAAAAGATAGGTTTGAGTTCAAATTTTGCATTGTGTGAGCTTCTACAGAT|\\$ TTTAGACAAGGACCGTTTTTACTAAGTAAAAGGGTGGAGAGGTTCCTGGGGTGGATTCCTAAGCAGTGCTTGTAA ACCATCGCGTGCAATGAGCCAGATGGAGTACCATGAGGGTTGTTATTTGTTGTTTTTAACAACTAATCAAGAGTG AGTGAACAACTATTTATAAACTAGATCTCCTATTTTTCAGAATGCTCTTCTACGTATAAATATGAAATGATAAAG ATGTCAAATATCTCAGAGGCTATAGCTGGGAACCCGACTGTGAAAGTATGTGATATCTGAACACATACTAGAAAG CTCTGCATGTGTTGTCCTTCAGCATAATTCGGAAGGGAAAACAGTCGATCAAGGGATGTATTGGAACATGTCG TGGCATCCCCTTTATTAATTCATTAAATTTCTGGATTTGGGTTGTGACCCAGGGTGCATTAACTTAAAAGATTCA CTAAAGCAGCACATAGCACTGGGAACTCTGGCTCCGAAAAACTTTGTTATATATCAAGGATGTTCTGGCTTTA CATTTTATTTATTAGCTGTAAATACATGTGTGGATGTGTAAATGGAGCTTGTACATATTGGAAAGGTCATTGTGG CTATCTGCATTTATAAATGTGTGGTGCTAACTGTATGTGTCTTTATCAGTGATGGTCTCACAGAGCCAACTCACT CTTATGAAATGGGCTTTAACAAAACAAGAAAGAAACGTACTTAACTGTGTGAAGAAATGGAATCAGCTTTTAATA AAATTGACAACATTTTATTACCAC

852/2825 FIGURE 746

MESKALLVLTLAVWLQSLTASRGGVAAADQRRDFIDIESKFALRTPEDTAEDTCHLIPGVAESVATCHFNHSSKT FMVIHGWTVTGMYESWVPKLVAALYKREPDSNVIVVDWLSRAQEHYPVSAGYTKLVGQDVARFINWMEEEFNYPL DNVHLLGYSLGAHAAGIAGSLTNKKVNRITGLDPAGPNFEYAEAPSRLSPDDADFVDVLHTFTRGSPGRSIGIQK PVGHVDIYPNGGTFQPGCNIGEAIRVIAERGLGDVDQLVKCSHERSIHLFIDSLLNEENPSKAYRCSSKEAFEKG LCLSCRKNRCNNLGYEINKVRAKRSSKMYLKTRSQMPYKVFHYQVKIHFSGTESETHTNQAFEISLYGTVAESEN IPFTLPEVSTNKTYSFLIYTEVDIGELLMLKLKWKSDSYFSWSDWWSSPGFAIQKIRVKAGETQKKVIFCSREKV SHLQKGKAPAVFVKCHDKSLNKKSG

853/2825 FIGURE 747

GGCGCCGACCCATGCTGGCTGGGAACGTGTCTCCCGGTGACGCAGCCCCGGGTGGGGAACGTGGTGCGGCGGCGG GACGGCAAACCCGGAGCTGCCGGCCGCGCGGGGAGGAGGACGCGGGTGCGGTCTAGGAAACGGAGCTGCGGGC GGAGGCTCCATGTTGGGAAGCGGCGCCGTTCGTGCTTGTTAGCGGGAATCCGGGAGCCGCGGGGTGAGCTGGCGG ACTACGAACTCGTCCAGAGGGTCGGCAGCGGCACCTACGGGGACGTCTATAAGGCCAGAAATGTACACACAGGAG AGCTGGCTGCAGTAAAAATCATTAAATTGGAGCCTGGAGATGATTTTTCTTTGATTCAACAAGAAATATTTATGG TTAAAGAATGTAAACATTGTAACATCGTTGCCTACTTTGGGAGTTATCTTAGTCGGGAAAAACTATGGATTTGTA TGGAATACTGTGGTGGCGGATCACTTCAAGATATTTACCATGTTACTGGACCATTATCAGAATTGCAAATAGCCT ATGTATGCAGAGAAACCTTACAGGGTCTTGCCTATTTGCATACTAAAGGCAAAATGCATAGAGATATCAAAGGTG CTAATATTTTATTGACAGACCATGGCGATGTAAAATTAGCTGACTTTGGTGTGGCTGCAAAAATAACAGCTACCA TTGCAAAACGAAAATCTTTCATTGGCACCCCTTACTGGATGGCCCCAGAAGTTGCAGCAGTAGAGAAGAATGGTG GCTACAACCAACTCTGTGATATCTGGGCAGTAGGAATAACAGCAATTGAACTTGGAGAACTTCAGCCACCTATGT TTGATCTCCACCCAATGAGGGCTCTCTTCTTAATGTCAAAAAGTAATTTTCAGCCTCCAAAACTAAAGGACAAAA AAAGACTTCTGACTCACACTTTTGTTGCACAGCCAGGTCTCTCTAGAGCCCTAGCAGTTGAACTGTTAGACAAAG TGAACAATCCAGATAACCACGCACATTACACTGAAGCAGATGACGATGACTTTGAGCCCCATGCAATCATTCGTC TTGAACCTCCTCTGAGAAAAGAAACAGAAGCACGAGATGAAATGGGATTGTCATCAGACCCAAATTTCATGTTAC AGTGGAATCCTTTTGTTGATGGTGCAAATACTGGCAAATCAACCTCAAAACGTGCAATACCACCTCCCCTACCTC CTAAGCCAAGGATAAGCAGTTACCCTGAAGACAACTTTCCGGATGAAGAAAAAGCATCAACCATAAAACATTGTC $\verb|CTGATTCAGAAAGCAGAGCTCCCCAAATTCTCAGAAGACAGAGTAGCCCAAGTTGTGGGCCTGTGGCAGAGACTT| \\$ CTTCTATTGGAAATGGTGATGGTATTTCAAAACTGATGAGTGAAAATACAGAAGGATCAGCACAAGCACCACAGT TACCACGAAAAAAGGACAAACGAGACTTCCCTAAACCAGCCATCAATGGCCTTCCACCCCACCCCAAAAGTTCTGA ATACAAAAGATCAGTACATTATTTTTGGAACTGAAGATGGTATTTACACACTGAATCTCAATGAGCTACATGAGG CAACGATGGAACAGTTATTTCCACGGAAGTGTACTTGGCTGTATGTTATCAATAATACTTTAATGTCATTATCAG CCCATATTCAAACTCACAGGTTTCCAGACCGAATACTACCAAGAAAATTCGCTTTAACAACAAAGATTCCTGATA CAAAAGGCTGCCACAAATGTTGCATAGTCAGAAACCCTTACACGGGACATAAATACCTCTGTGGAGCTTTACAGT CTGGAATTGTTTTACTTCAGTGGTATGAGCCAATGCAGAAATTCATGTTGATAAAGCACTTTGATTTTCCTTTGC CAAGTCCTTTGAATGTTTTTGAAATGCTGGTGATACCTGAACAGGAATACCCTATGGTCTGTGTAGCTATTAGCA ACAAATTTGTGAAAATTGTAAATCTACAAGGAAAATTAAAATCAAGTAAGAAACTGGCCTCTGAGTTAAGTTTTG ATTTTCGCATTGAATCTGTAGTATGCCTTCAAGACAGTGTGTTTGGCTTTCTGGAAACATGGGATGCAGGGTAAAA GCTTCAAGTCAGATGAGGTTACCCAGGAGATTTCAGATGAAACAAGAGTTTTCCGCTTATTAGGATCAGACAGGG TTGTCGTTTTGGAAAGTAGGCCAACAGAAAATCCTACTGCACACAGCAATCTCTACATCTTGGCTGGACATGAAA ATAGTTACTAAGCAACAGAAACTGATCTCAAATGACAGGAAAATGAATATACTCCATTGAAAGGAAAAATAAGGA AATTCAATACAAACTGCACTATGATTTGCTTTAACTATTATGGGTTATATTGCAAATGATCTGTACTTTAGGGTA

854/2825 FIGURE 748

MEAPLRPAADILRRNPQQDYELVQRVGSGTYGDVYKARNVHTGELAAVKIIKLEPGDDFSLIQQEIFMVKECKHC
NIVAYFGSYLSREKLWICMEYCGGGSLQDIYHVTGPLSELQIAYVCRETLQGLAYLHTKGKMHRDIKGANILLTD
HGDVKLADFGVAAKITATIAKRKSFIGTPYWMAPEVAAVEKNGGYNQLCDIWAVGITAIELGELQPPMFDLHPMR
ALFLMSKSNFQPPKLKDKTKWSSTFHNFVKIALTKNPKKRPTAERLLTHTFVAQPGLSRALAVELLDKVNNPDNH
AHYTEADDDDFEPHAIIRHTIRSTNRNARAERTASEINFDKLQFEPPLRKETEARDEMGLSSDPNFMLQWNPFVD
GANTGKSTSKRAIPPPLPPKPRISSYPEDNFPDEEKASTIKHCPDSESRAPQILRRQSSPSCGPVAETSSIGNGD
GISKLMSENTEGSAQAPQLPRKKDKRDFPKPAINGLPPTPKVLMGACFSKVFDGCPLKINCATSWIHPDTKDQYI
IFGTEDGIYTLNLNELHEATMEQLFPRKCTWLYVINNTLMSLSEGKTFQLYSHNLIALFEHAKKPGLAAHIQTHR
FPDRILPRKFALTTKIPDTKGCHKCCIVRNPYTGHKYLCGALQSGIVLLQWYEPMQKFMLIKHFDFPLPSPLNVF
EMLVIPEQEYPMVCVAISKGTESNQVVQFETINLNSASSWFTEIGAGSQQLDSIHVTQLERDTVLVCLDKFVKIV
NLQGKLKSSKKLASELSFDFRIESVVCLQDSVLAFWKHGMQGKSFKSDEVTQEISDETRVFRLLGSDRVVVLESR
PTENPTAHSNLYILAGHENSY

855/2825 FIGURE 749

856/2825 FIGURE **750**

 ${\tt MAGKAHRLSAEERDQLLPNLRAVGWNELEGRDAIFKQFHFKDFNRAFGFMTRVALQAEKLDHHPEWFNVYNKVHI}\\ {\tt TLSTHECAGLSERDINLASFIEQVAVSMT}$

.857/2825 FIGURE **751**

TGCCGCAGCCCCGCCCGCAGAGCTTTTGAAAGGCGGCGGAGGCGGCGAGCGCCA**TG**GCCAGTCCGGGCT GCCTGCTGTGCGTGCTGGGCCTACTCTGCGGGGCGGCGAGCCTCGAGCTGTCTAGACCCCACGGCGACACCG CCAAGAAGCCCATCATCGGAATATTAATGCAAAAATGCCGTAATAAAGTCATGAAAAACTATGGAAGATACTATA TTGCTGCGTCCTATGTAAAGTACTTGGAGTCTGCAGGTGCGAGAGTTGTACCAGTAAGGCTGGATCTTACAGAGA AAGACTATGAAATACTTTTCAAATCTATTAATGGAATCCTTTTCCCTGGAGGAAGTGTTGACCTCAGACGCTCAG ATTATGCTAAAGTGGCCAAAATATTTTATAACTTGTCCATACAGAGTTTTGATGATGAGAGACTATTTTCCTGTGT GGGGCACATGCCTTGGATTTGAAGAGCTTTCACTGCTGATTAGTGGAGAGTGCTTATTAACTGCCACAGATACTG TTGACGTGGCAATGCCGCTGAACTTCACTGGAGGTCAATTGCACAGCAGAATGTTCCAGAATTTTCCTACTGAGT TGTTGCTGTCATTAGCAGTAGAACCTCTGACTGCCAATTTCCATAAGTGGAGCCTCTCCGTGAAGAATTTTACAA TGAATGAAAAGTTAAAGAAGTTTTTCAATGTCTTAACTACAAATACAGATGGCAAGATTGAGTTTATTTCAACAA TGGAAGGATATAAGTATCCAGTATATGGTGTCCAGTGGCATCCAGAGAAAGCACCTTATGAGTGGAAGAATTTGG AAAACAACCATCATTTTAAATCTGAATCTGAAGAGGAGAAAGCATTGATTTATCAGTTCAGTCCAATTTATACTG GAAATATTTCTTCATTTCAGCAATGTTACATATTTGAT**TGA**AAGTCTTCAATTTGTTAACAGAGCAAATTTGAAT AATTCCATGATTAAACTGTTAGAATAACTTGCTACTCATGGCAAGATTAGGAAGTCACAGATTCTTTTCTATAAT GTGCCTGGCTCTGATTCTTCATTATGTATGTGACTATTTATATAACATTAGATAATTAAATAGTGAGACATAAAT AAAAA

858/2825 FIGURE **752**

MASPGCLLCVLGLLLCGAASLELSRPHGDTAKKPIIGILMQKCRNKVMKNYGRYYIAASYVKYLESAGARVVPVR LDLTEKDYEILFKSINGILFPGGSVDLRRSDYAKVAKIFYNLSIQSFDDGDYFPVWGTCLGFEELSLLISGECLL TATDTVDVAMPLNFTGGQLHSRMFQNFPTELLLSLAVEPLTANFHKWSLSVKNFTMNEKLKKFFNVLTTNTDGKI EFISTMEGYKYPVYGVQWHPEKAPYEWKNLDGISHAPNAVKTAFYLAEFFVNEARKNNHHFKSESEEEKALIYQF SPIYTGNISSFQQCYIFD

859/2825 FIGURE **753**

GAATTCCAACT**ATG**GAGACCCAAATGTCTCAGAATGTATGTCCCAGAAACCTGTGGCTGCTTCAACCATTGACAG TTTTGCTGCTGCTTCTGCAGACAGTCAAGCTGCAGCTCCCCCAAAGGCTGTGCTGAAACTTGAGCCCCCGT GGATCAACGTGCTCCAGGAGGACTCTGTGACTCTGACATGCCAGGGGGGCTCGCAGCCCTGAGAGCGACTCCATTC AGTGGTTCCACAATGGGAATCTCATTCCCACCCACACGCAGCCCAGCTACAGGTTCAAGGCCAACAACAATGACA GCGGGGAGTACACGTGCCAGACTGGCCAGACCAGCCTCAGCGACCCTGTGCATCTGACTGTGCTTTCCGAATGGC TGGTGCTCCAGACCCCTCACCTGGAGTTCCAGGAGGGGAGAAACCATCATGCTGAGGTGCCACAGCTGGAAGGACA AGCCTCTGGTCAAGGTCACATTCTTCCAGAATGGAAAATCCCAGAAATTCTCCCGTTTGGATCCCACCTTCTCCA TCCCACAAGCAAACCACAGTCACAGTGGTGATTACCACTGCACAGGAAACATAGGCTACACGCTGTTCTCATCCA AGCCTGTGACCATCACTGTCCAAGTGCCCAGCATGGGCAGCTCTTCACCAATGGGGATCATTGTGGCTGTGGTCA TTGCGACTGCTGTAGCAGCCATTGTTGCTGCTGTAGTGGCCTTGATCTACTGCAGGAAAAAGCGGATTTCAGCCA ATTCCACTGATCCTGTGAAGGCTGCCCAATTTGAGCCACCTGGACGTCAAATGATTGCCATCAGAAAGAGACAAC TTGAAGAAACCAACAATGACTATGAAACAGCTGACGGCGGCTACATGACTCTGAACCCCAGGGCACCTACTGACG ATGATAAAAACATCTACCTGACTCTTCCTCCCAACGACCATGTCAACAGTAATAACTAAAGAGTAACGTTATGCC ATGTGGTCATACTCTCAGCTTGCTGAGTGGATGACAAAAAGGGGGAATTGTTAAAGGAAAATTTAAATGGAGAC TGGAAAAATCCTGAGCAAACAAAACCACCTGGCCCTTAGAAATAGCTTTAACTTTGCTTAAACTACAAACACAAG CAAAACTTCACGGGGTCATACTACATACAAGCATAAGCAAAACTTAACTTGGATCATTTCTGGTAAATGCTTATG TTAGAAATAAGACAACCCCAGCCAATCACAAGCAGCCTACTAACATATAATTAGGTGACTAGGGACTTTCTAAGA AATAAATACTTGCCTGTGACATTTTGCCACTGGAACACTAAACTTCATGAATTGCGCCTCAGATTTTTCCTTTAA CATCTTTTTTTTTTTGACAGAGTCTCAATCTGTTACCCAGGCTGGAGTGCAGTGGTGCTATCTTGGCTCACTGC AAACCCGCCTCCCAGGTTTAAGCGATTCTCATGCCTCAGCCTCCCAGTAGCTGGGATTAGAGGCATGTGCCATCA CTCGAACTTCTGGCCTCTAGCGATCTGCCCGCCTCGGCCTCCCAAAGTGCTGGGATGACCAGCATCAGCCCCAAT GTCCAGCCTCTTTAACATCTTCTTTCCTATGCCCTCTCTGTGGATCCCTACTGCTGGTTTCTGCCTTCTCCATGC TGAGAACAAATCACCTATTCACTGCTTATGCAGTCGGAAGCTCCAGAAGAACAAAGAGCCCAATTACCAGAACC ACATTAAGTCTCCATTGTTTTTGCCTTGGGATTTGAGAAGAGAATTAGAGAGGTGAGGATCTGGTATTTCCTGGAC TAAATTCCCCTTGGGGAAGACGAAGGGATGCTGCAGTTCCAAAAGAGAGGACTCTTCCAGAGTCATCTACCTGA CGTTCTTGACATCAAGAATCTTCTGTTCCACATCCACACACCCAATACAATTAGTCAAACCACTGTTATTAACAG ATGTAGCAACATGAAAGACGCTATGTTACAGGTTACATGAGAGCAATCATGTAAGTCTATATGACTTCAGAAATG TTAAAATAGACTAACCTCTAACAACAAATTAAAAGTGCGGAATTC

860/2825 FIGURE 754

METQMSQNVCPRNLWLLQPLTVLLLLASADSQAAAPPKAVLKLEPPWINVLQEDSVTLTCQGARSPESDSIQWFH NGNLIPTHTQPSYRFKANNNDSGEYTCQTGQTSLSDPVHLTVLSEWLVLQTPHLEFQEGETIMLRCHSWKDKPLV KVTFFQNGKSQKFSRLDPTFSIPQANHSHSGDYHCTGNIGYTLFSSKPVTITVQVPSMGSSSPMGIIVAVVIATA VAAIVAAVVALIYCRKKRISANSTDPVKAAQFEPPGRQMIAIRKRQLEETNNDYETADGGYMTLNPRAPTDDDKN IYLTLPPNDHVNSNN

861/2825 FIGURE 755

862/2825 FIGURE 756

MAKVSELYDVTWEEMRDKMRKWREENSRNSEQIVEVGEELINEYASKLGDDIWIIYEQVMIAALDYGRDDLALFC LQELRRQFPGSHRVKRLTGMRFEAMERYDDAIQLYDRILQEDPTNTAARKRKIAIRKAQGKNVEAIRELNEYLEQ FVGDQEAWHELAELYINEHDYAKAAFCLEELMMTNPHNHLYCQQYAEVKYTQGGLENLELSRKYFAQALKLNNRN MRALFGLYMSASHIASNPKASAKTKKDNMKYASWAASQINRAYQFAGRSKKETKYSLKAVEDMLETLQITQS

863/2825 FIGURE 757

TGAACATCAAGTTGGTGCT**ATG**GCAAGGCTGGGAAACTGCAGCCTGACTTGGGCTGCCCTGATCATCCTGCTGCT CCCCGGAAGTCTGGAGGAGTGCGGGCACATCAGTGTCTCAGCCCCCATCGTCCACCTGGGGGGATCCCATCACAGC CTCCTGCATCATCAAGCAGAACTGCAGCCATCTGGACCCGGAGCCACAGATTCTGTGGAGACTGGGAGCAGAGCT TCAGCCCGGGGGCAGCAGCAGCTCTGTCTGATGGGACCCAGGAATCTATCATCACCCTGCCCCACCTCAACCA CACTCAGGCCTTTCTCTCCTGCTGCCTGAACTGGGGCAACAGCCTGCAGATCCTGGACCAGGTTGAGCTGCGCGC AGGCTACCCTCCAGCCATACCCCACAACCTCTCCTGCCTCATGAACCTCACAACCAGCAGCCTCATCTGCCAGTG CCAAGGGGACTCCATCCTGGACTGCCTGCCCAAGGACGGCAGAGCCACTGCTGCATCCCACGCAAACACCTGCT GTTGTACCAGAATATGGGCATCTGGGTGCAGGCAGAATGCGCTGGGGACCAGCATGTCCCCACAACTGTGTCT TGATCCCATGGATGTTGTGAAACTGGAGCCCCCATGCTGCGGACCATGGACCCCAGCCCTGAAGCGGCCCCTCC CCAGGCAGGCTGCCTACAGCTGTGCTGGGAGCCATGGCAGCCTGCACATAAATCAGAAGTGTGAGCTGCG CCACAAGCCGCAGCGTGGAGAAGCCAGCTGGGCACTGGTGGGCCCCCTCCCCTTGGAGGCCCTTCAGTATGAGCT CGACTGGAGCCCCAGCCTGGAGCTGAGAACTACCGAACGGGCCCCCACTGTCAGACTGGACACATGGTGGCGGCA AGGTTATGTGGTTTCTTGGAGACCCTCAGGCCAGGCTGGGGCCATCCTGCCACTCTGCAACACCACAGAGCTCAG CTGCACCTTCCACCTGCCTTCAGAAGCCCAGGAGGTGGCCCTTGTGGCCTATAACTCAGCCGGGACCTCTCGCCC CACCCCGGTGGTCTTCTCAGAAAGCAGAGGCCCAGCTCTGACCAGACTCCATGCCATGGCCCGAGACCCTCACAG CCTCTGGGTAGGCTGGGAGCCCCCAATCCATGGCCTCAGGGCTATGTGATTGAGTGGGGCCTGGGCCCCCCAG CGCGAGCAATAGCAACAAGACCTGGAGGATGGAACAGAATGGGAGAGCCACGGGGTTTCTGCTGAAGGAGAACAT CAGGCCCTTTCAGCTCTATGAGATCATCGTGACTCCCTTGTACCAGGACACCATGGGACCCTCCCAGCATGTCTA TGCCTACTCTCAAGAAATGGCTCCCTCCCATGCCCCAGAGCTGCATCTAAAGCACATTGGCAAGACCTGGGCACA GCTGGAGTGGCTGAGCCCCTGAGCTGGGGAAGAGCCCCCTTACCCACTACACCATCTTCTGGACCAACGC TCAGAACCAGTCCTTCTCCGCCATCCTGAATGCCTCCTCCCGTGGCTTTGTCCTCCATGGCCTGGAGCCCGCCAG TCTGTATCACATCCACCTCATGGCTGCCAGCCAGGCTGGGGCCACCAACAGTACAGTCCTCACCCTGATGACCTT GACCCCAGAGGGGTCGGAGCTACACATCATCCTGGGCCTGTTCGGCCTCCTGCTGTTGCTCACCTGCCTCTGTGG AACTGCCTGGCTCTGTTGCAGCCCCAACAGGAAGAATCCCCTCTGGCCAAGTGTCCCAGACCCAGCTCACAGCAG CCTGGGCTCCTGGGTGCCCACATCATGGAGGAGGATGCCTTCCAGCTGCCCGGCCTTGGCACGCCACCCATCAC CAAGCTCACAGTGCTGGAGGAGGATGAAAAGAAGCCGGTGCCCTGGGAGTCCCATAACAGCTCAGAGACCTGTGG CCTCCCCACTCTGGTCCAGACCTATGTGCTCCAGGGGGGACCCAAGAGCAGTTTCCACCCAGCCCCAATCCCAGTC TGGCACCAGCGATCAGGTCCTTTATGGGCAGCTGCTGGGCAGCCCCACAAGCCCAGGGCCAGGGCACTATCTCCG CTGTGACTCCACTCAGCCCCTCTTGGCGGGCCTCACCCCCAGCCCCAAGTCCTATGAGAACCTCTGGTTCCAGGC ${\tt CAGCCCTTGGGGACCCTGGTAACCCCAGCCCCAAGCCAGGGAGGACGACTGTGTCTTTGGGCCACTGCTCAACTT}$ CCCCCTCCTGCAGGGGATCCGGGTCCATGGGATGGAGGCGCTGGGGAGCTTC**TAG**GGCTTCCTGGGGTTCCCTTC TTGGGCCTGCTCTTAAAGGCCTGAGCTAGCTGGAGAAGAGGGGAGGGTCCATAAGCCCATGACTAAAAACTACC CCAGCCCAGGCTCTCACCATCTCCAGTCACCAGCATCTCCCTCTCCCCAATCTCCATAGGCTGGGCCTCCCAG AACTTCAGTATTGTAAAC

864/2825 FIGURE **758**

MARLGNCSLTWAALIILLLPGSLEECGHISVSAPIVHLGDPITASCIIKQNCSHLDPEPQILWRLGAELQPGGRQ
QRLSDGTQESIITLPHLNHTQAFLSCCLNWGNSLQILDQVELRAGYPPAIPHNLSCLMNLTTSSLICQWEPGPET
HLPTSFTLKSFKSRGNCQTQGDSILDCVPKDGQSHCCIPRKHLLLYQNMGIWVQAENALGTSMSPQLCLDPMDVV
KLEPPMLRTMDPSPEAAPPQAGCLQLCWEPWQPGLHINQKCELRHKPQRGEASWALVGPLPLEALQYELCGLLPA
TAYTLQIRCIRWPLPGHWSDWSPSLELRTTERAPTVRLDTWWRQRQLDPRTVQLFWKPVPLEEDSGRIQGYVVSW
RPSGQAGAILPLCNTTELSCTFHLPSEAQEVALVAYNSAGTSRPTPVVFSESRGPALTRLHAMARDPHSLWVGWE
PPNPWPQGYVIEWGLGPPSASNSNKTWRMEQNGRATGFLLKENIRPFQLYEIIVTPLYQDTMGPSQHVYAYSQEM
APSHAPELHLKHIGKTWAQLEWVPEPPELGKSPLTHYTIFWTNAQNQSFSAILNASSRGFVLHGLEPASLYHIHL
MAASQAGATNSTVLTLMTLTPEGSELHIILGLFGLLLLLTCLCGTAWLCCSPNRKNPLWPSVPDPAHSSLGSWVP
TIMEEDAFQLPGLGTPPITKLTVLEEDEKKPVPWESHNSSETCGLPTLVQTYVLQGDPRAVSTQPQSQSGTSDQV
LYGQLLGSPTSPGPGHYLRCDSTQPLLAGLTPSPKSYENLWFQASPLGTLVTPAPSQEDDCVFGPLLNFPLLQGI
RVHGMEALGSF

865/2825 FIGURE 759

GCTGACTCCAGTGTCCCGAGAGGCGCCGCTTCTTCCGCTTTCTCGTCAGGCTCCTGCAACCCCAGGCATGAACCA AGGTTTCTGAACTACTGGGCGGAGCCAACGTCTCTTCTTCTCCCGCTCTGGCGGAGGCTTTGTCGCTGCGGGC TGGGCCCCAGGGTGTCCCCCATCGCGGGGCCGCGGGTGGAGGTCGATGGCAGCATCATGGAAGGGGGGCGGCCAGA GCACGCCAGGCCTGAGGCCTCAACATTTATCTGGACTGGAAATGATTCGAGATTTGTGTGATGGGCAACTGGAGG GGGCAGAAATTGGCTCAACAGAAATAACCTTTACACCAGAGAAGATCAAAGGTGGAATCCACACAGCAGATACCA AGACAGCAGGGAGTGTGTGCCTCTTGATGCAGGTCTCAATGCCGTGTGTTCTCTTTGCTGCTTCTCCATCAGAAC TTCATTTGAAAGGTGGAACTAATGCTGAAATGGCACCACAGATCGATTATACAGTGATGGTCTTCAAGCCAATTG TTGAAAAATTTGGTTTCATATTTAATTGTGACATTAAAACAAGGGGATATTACCCAAAAGGGGGTGGTGAAGTGA TTGTTCGAATGTCACCAGTTAAACAATTGAACCCTATAAATTTAACTGAGCGTGGCTGTGTGACTAAGATATATG GAAGAGCTTTCGTTGCTGGTGTTTTTGCCATTTAAAGTAGCAAAAGATATGGCAGCGGCAGCAGTTAGATGCATCA GAAAGGAGATCCGGGATTTGTATGTTAACATCCAGCCTGTTCAAGAACCTAAAGACCAAGCATTTGGCAATGGAA ATGCAGACAAAGTTGGAATTGAAGCTGCCGAAATGCTATTAGCAAATCTTAGACATGGTGGTACTGTGGATGAGT ATCTGCAAGACCAGCTGATTGTTTTCATGGCATTAGCCAATGGAGTTTCCAGAATAAAAACAGGACCAGTTACAC TCCATACGCAAACCGCGATACATTTTGCTGAACAAATAGCAAAGGCTAAATTTATTGTGAAGAAATCAGAAGATG AAGAAGACGCCGCTAAAGATACTTATTATTGAATGCCAAGGAATTGGGATGACAAATCCAAATCTA**TAG**AGTA TTTGCCTCTTAAATGATACCTCATTGATATATTGCACTATTTCATAAATACTATAAAATAATGACTAGGAAGTAA CTTATTAAAGGCTATGACTTAAATTTGAAGATGAAGTACAGTGTTCTAGGTTTGCTGAGAAGGCTTCATTAAATT AATCTCACTTTGAATATCTCCTGAGAGATGGACAATGAAATATCAGTTGGTGGATATGTGTGATAGCTGATTTCA ATATTGAAGTATTGAAATAAAATATTCTTTACACCTGAG

866/2825 FIGURE 760

MAGPRVEVDGSIMEGGGQILRVSTALSCLLGLPLRVQKIRAGRSTPGLRPQHLSGLEMIRDLCDGQLEGAEIGST EITFTPEKIKGGIHTADTKTAGSVCLLMQVSMPCVLFAASPSELHLKGGTNAEMAPQIDYTVMVFKPIVEKFGFI FNCDIKTRGYYPKGGGEVIVRMSPVKQLNPINLTERGCVTKIYGRAFVAGVLPFKVAKDMAAAAVRCIRKEIRDL YVNIQPVQEPKDQAFGNGNGIIIIAETSTGCLFAGSSLGKRGVNADKVGIEAAEMLLANLRHGGTVDEYLQDQLI VFMALANGVSRIKTGPVTLHTQTAIHFAEQIAKAKFIVKKSEDEEDAAKDTYIIECQGIGMTNPNL

867/2825 FIGURE **761A**

GCGGCCGCTGGCTCCCGGCCGCCACCTTCCCCGTGCAAGACCCGGCCAGCGGCGCCGCTCTGGGCATGGTAG CCGCCAAGGAGGAGGTCATTACTTCGGAAGTGGTACAATTTAATGATACAAAATAAGGATGACCTTGCCAGAA TAATCACAGCTGAAAGTGGAAAGCCACTGAAGGAGGCACATGGAGAAATTCTCTATTCCGCCTTTTTCCTAGAGT GGTTCTCTGAGGAAGCCCGCCGTGTTTACGGAGACATTATCTACACCCCGGCAAAGGACAGGCGGGGCCCTGGTC CTCAAGCAGCCCATAGGCGTGGGCTGCAGTCATCACCCCGTGGAATTTCCCCAGTGCCATGATCACCCGGAAGGT GGGGGCCGCCTGGCAGCCGCTGTACTGTCGTGGTGAAGCCTGCCGAAGACACGCCCTTCTCCGCCCTGGCCCT GGCTGAGCTTGCAAGCCAGGCTGGGATTCCTTCAGGTGTATACAATGTTATTCCCTGTTCTCGAAAGAATGCCAA GGAAGTAGGGGAGCAATTTGTACTGATCCTCTGGTGTCCAAAATTTCCTTTACTGGTTCAACAACTACAGGAAA GATCCTGTTGCACCACGCAGCAAACTCTGTGAAAAGGGTCTCTATGGAGCTGGGCGGCCTTGCTCCATTTATAGT ATTTGACAGTGCCAACGTGGACCAGGCTGTAGCAGGGCCATGGCATCTAAATTTAGGAACACTGGACAGACTTG TGTTTGCTCAAACCAATTCTTGGTGCAAAGGGGCATCCATGATGCCTTTGTAAAAGCATTCGCCGAGGCCATGAA AAAGGTGGAGAAACAGGTGAATGATGCCGTTTCTAAAGGTGCCACCGTTGTGACAGGTGGAAAACGACACCAACT TGGAAAAAATTTCTTTGAGCCTACCCTGCTGTGCAATGTCACCCAGGACATGCTGTGCACTCATGAAGAGACTTT CGGGCCTCTGGCACCAGTTATCAAGTTCGATACAGAGGAGGAGGCTATAGCAATCGCTAACGCAGCTGATGTTGG GTTAGCAGGTTATTTTTACTCTCAAGACCCAGCCCAGATCTGGAGAGTGGCAGAGCAGCTGGAAGTGGGCATGGT TGGCGTCAACGAAGGATTAATTTCCTCTGTGGAGTGCCCTTTTGGTGGAGTGAAGCAGTCCGGCCTTGGGCGAGA GGGGTCCAAGTATGGCATTGATGAGTATCTGGAACTCAAGTATGTGTGTTACGGGGGGCTTG**TAG**GATTCTTTGGT GGTTTTCAGAATTATGAATTTTTCAGAACTCATCCAGTCCTTGTAATCTTAAACAGATGCAAATCCTACCCCTGC CCTTAATGTAACTAGGGACCAATATGTGCCACGTGCCTGTGGCTGCAGACTCCCAGAGAACCAGCACTGGGTTTA ATGGGCCGTCACAAAGGCTTGATCACTGCCTGGGCAGTGGCACAACCATCCCCCATGTCGCTACAGAACATGGGC GCCAGCTCAGCACAGAAGACACCGAATGTGCATCTGGAGGTGTTGCGGCAGAGGTTTGAGTGAACCCTCTTTCAA AGACAATAAATAGCACAGAATTGTCCGTGCTTCTGTGAGGCACGAAGGAGCACCTCTCCCTATTCCTGAACCATA TTTTAATTGATGCCTTTATTGTCCCAAATTTACCACTTAAAGAATTCTATTTTCATTTCTGTCCTGTTTGCCACT TCATTTCATTTCCTTGAGTAAATGAGCTGAAGTGTAAAGCAAAGAATAGCAGAAGCAGCTTGGTAGGATGGAAGG CACGCAGTGGCATGGCCTGCCTCCCTGAGTGGCCTTGTTGCCTTGGGCGTGTTTCAGCACCTCCCATCCC ATGGATACAGTGGAGATTCCATACAGGCCTTACTTAGCTCCTTGCGATACTGTGAGCCAGAGAGAAAGTAAAAGC CCTACTGTCAGTGGCTTTGCCAAGATGGGAGGGCCAGTGCCCACTTGCCCACCGACCTGAGCCTGAGTAAGTG ACCACAGTCAACCCAAAAGGGAAAGAATTTCTGTAATTGCCAACCCACTTTGTAGAAGAGGCATGAGCCCTTGAA TTTCAGGCACCTTTTATCTAATGAGAGTTTTTACAGCTCATCATTCAAATAACGCACAAAAAAACAAAAGCCGCTA TGACCGGCGTTGCCTCTCACTTTGAAAGAAAGACTCCGTAATCTTAAACAGATGCAAATCCAATTAGGCGGCCAA GTAGGTGGAGACCACCTTCCTTTCCCCATCTCTGTCTGTAGACTCCACTTACATCTTGTGGTTATTCTG GGTTGTGGAAGGAAAATGTGAATGAAAGTAGTGCTGGATGCAATTACTCCTGAACCCCACCTAGGATTTAACATC CCTCATTGAGCCACTGGGAAGCTATGGTTGAGTAACTAAAGGTCTGGCAAGACCCTGGGCACACAGGGAGCTGCC GTGACAGTGCCCCGGCCCTGTCAAGAATCACATACTCAGCATTTTTGACAGTTAATGTCTTCTCTTTATTTGTGTA NNNNNNNTGCTCCAACATTATTTTTTATAACTCTGGATTACCGATGCTGAGATGTGTGCAGGAATACTTCCAA

868/2825 FIGURE 761B

ATAAATTTTTAACATTGTAAGCCAGTAGATTCTTTGGCTGGGTGGCAAATTTTGAGTCTTCTTGCATATTTGGCT CATGGGTTCATAAATTGAGGCCAGAGACTCATTGTATATCCTTGATCAATTTTCTAATTAAAGGAAATAGCTTAT TGTTTAGTAAAATGGCCCTTGGTTGTGGGAAGAAAGAGAATTATTTTTAACTGAACGTTTTTGTTGTTTTTTAAA TAAAGTAGCCATATGTAACGGGATTTTAAAAATATATACATTTTACAACATGGACTAGAAAAGGACCATTAGATC AGGGGTTGGCAAACTATGGCCCACGAATCAAACCTGGCCTGCCACCTGTTTTTATACAACCTGTGAGCTAAGAAT GGTTTATATGTTTTCAAATGATTGGGCAGAAAATCAAAAGAAGAACAATATTTCATGACACCTGAAAATTCTATG GGACTGCTTTCTTGGCAGAGTTGAATAGTTGGGACAGGGACAGTATAGCCTGCAAAGCCTAAAAGATTTACTACT ${\tt TGTCTCCTTAAAGAAAAGTTTGCCAGCTCCTGATCTAGACCAAGAGGGAGATATTGACCTTACTTGCATTTATC}$ CGGCATTATTTCAACACCTCCTGTGAAGAAAGTACTTTTCTGGCATAGCTACCGTGAGTTCTCAGCAGGAACTTC CCAGCGATGATAATGTACATGAAATTAGCCACGTAACAAGAGTCTATTTGCCTTGAAAAAACAATTATCCTTGAA CTTCCTCCAGCTATGCAGGGATTCTCATTAAAGAGCTGATCTGCATCTGTTTATCCATTTTAGTAGCTTTTAAAT ATGATTTTATGTGAAGTGCCATTTCTTTATTCTACTAAAGAGAGACAGCAGAAATAAAGATCAGAAGCAAATGTG AGCAAATCTGTTTACTGTTAACTTCAATTATGAACTTGAAATTGTGCCACATGACTTTTTCCTCTAAAAACTGG TTTCATGCTATACTTTGTGGGATAAAACTTGGGAATGAGTGTGGTAAGAAATTTATAAAGTTTTGCTTTTAAAAC GTGGACATAACTCATTTTTCTAGTTTTTGACAATTGTGTGTTTTTAGTGTCTAGTCTGCAGAGAGCTGTGTGATTA

869/2825 FIGURE 762

GRWLPAAATFPVQDPASGAALGMVADCGVREARAAVRAAYEAFCRWREVSAKERSSLLRKWYNLMIQNKDDLARI
ITAESGKPLKEAHGEILYSAFFLEWFSEEARRVYGDIIYTPAKDRRGPGPQAAHRAWAAVITPWNFPSAMITRKV
GAALAAGCTVVVKPAEDTPFSALALAELASQAGIPSGVYNVIPCSRKNAKEVGEAICTDPLVSKISFTGSTTTGK
ILLHHAANSVKRVSMELGGLAPFIVFDSANVDQAVAGAMASKFRNTGQTCVCSNQFLVQRGIHDAFVKAFAEAMK
KNLRVGNGFEEGTTQGPLINEKAVEKVEKQVNDAVSKGATVVTGGKRHQLGKNFFEPTLLCNVTQDMLCTHEETF
GPLAPVIKFDTEEEAIAIANAADVGLAGYFYSQDPAQIWRVAEQLEVGMVGVNEGLISSVECPFGGVKQSGLGRE
GSKYGIDEYLELKYVCYGGL

870/2825 FIGURE 763

871/2825 FIGURE 764

MTGYTPDEKLRLQQLRELRRRWLKDQELSPREPVLPPQKMGPMEKFWNKFLENKSPWRKMVHGVYKKSIFVFTHV LVPVWIIHYYMKYHVSEKPYGIVEKKSRIFPGDTILETGEVIPPMKEFPDQHH

872/2825 FIGURE 765

TTTCCACATGTGAGGGGCTGAGGGGCTCAAGGAGGGGAGCATCTCCACTGGGTGGAGGCTGGGGGTCCCAGCAGG AAATGGTGAGACAAAGGGCGCTGGCTGGCAGGGAGACAGCACAGGAAGGTCCTAGAGCTTCCTCAGTGCAGCTGG ACTCTCCTGGAGACCTTCACACACCCTGATATCTGGGCCTTGCCCGACGAGGGTGCTTTCACTGGTCTGCACCAT TAGGCCTCAGTGACTTCCCTGAAAGCAAGAATTCCACTTTCTTGCTGTTGTGATGGTGGTAAGGGAACGGGCCTG TCAGCACCTACCCCAAGTCCGGCACCACCTGGGTGAGCCAGATTCTGGACATGATCTACCAGGGTGGTGACCTGG TGGAGACTCTGAAAGACACCCGGCCCCACGACTCCTGAAGACACCTGCCCCTGGCTCTGCTCCCCAGACTC TGTTGGATCAGAAGGTCAAGGTGGTCTATGTTGCCCGCAACGCAAAGGATGTGGCAGTTTCCTACTACCACTTCT ACCACATGGCCAAGGTGCACCCTGAGCCTGGGACCTGGGACAGCTTCCTGGAGAAGTTCATGGTCGGAGAAGTGT ATGAAGACATGAAGGAGAACCCCAAAAGGGAGATTCAAAAGATCCTGGAGTTTGTGGGGCACTCCCTGCCAGAGG AGACCGTGGACTTCATGGTTCAGCACACGTCGTTCAAGGAGATGAAGAAGAACCCTATGACCAACTACACCACCG TCCCCCAGGAGTTCATGGACCACAGCATCTCCCCCTTCATGAGGAAAGGCATGGCTGGGGACTGGAAGACCACCT TCACCGTGGCGCAGAATGAGCGCTTCGATGCGGACTATGCGGAGAAGATGGCAGGCTGCAGCCTCAGCTTCCGCT CTGAGCTG**TGA**GAGGGGCTCCTGGGGTCACTGCAGAGGGGAGTGTGCGAATCAAACCTGACCAAGCGGCTCAAGAA TAAAATATGAATTGAGGGCCTGGGACGGTAGGTCATGTCTGTAATCCCAGCAATTTGGAGGCTGAGGTGGGAGGA ACCAATTTTT

873/2825 FIGURE 766

MELIQDTSRPPLEYVKGVPLIKYFAEALGPLQSFQARPDDLLISTYPKSGTTWVSQILDMIYQGGDLEKCHRAPI FMRVPFLEFKAPGIPSGMETLKDTPAPRLLKTHLPLALLPQTLLDQKVKVVYVARNAKDVAVSYYHFYHMAKVHP EPGTWDSFLEKFMVGEVSYGSWYQHVQEWWELSRTHPVLYLFYEDMKENPKREIQKILEFVGHSLPEETVDFMVQ HTSFKEMKKNPMTNYTTVPQEFMDHSISPFMRKGMAGDWKTTFTVAQNERFDADYAEKMAGCSLSFRSEL

874/2825 FIGURE 767

GACGGCGGTGGCGCGTCGGAGGCGCCTCCGGGGGACGGTGGCGGCTCCCGGCGGTGAGGCCGCCCTGTCCGGG GATCGTCGAGGGACGGGGGGGCTTGGGCCAGCGGCGGCGGCGCCTGGGACGCAGGCCGCGCAGGCC CAAGGTCCCGGAGGCT<u>ATC</u>GCAGCGGCTACCATCGTGCACGACACGTCTGAGGCCGTGGAGCTCTGCCCTGCTTA CGGCTTGTACCTGAAGCCCATCACCAAGATGACCATCAGCGTGGCACTCCCGCAGCTGAAGCAGCCGGGGAAGTC CATCTCCAACTGGGAGGTGATGGAGGGCTGAAGGGCATGGTGCAGAACCACCAGTTCTCCACGCTGCGTATTTC CCTGGACGCAAGACCATCAAGCTCAGCGGCTTCTCCGACATCCTGAAGGTGCGCGGCCGAGTTCAAGATCGA CTTCCCCACCCGCCACGACTGGGACTCCTTCTTCCGCGACGCCAAGGACATGAACGAGACCCTGCCGGGGGAGCG GCCGGACACCATCCACCTGGAGGGGCTGCCCTGCAAGTGGTTCGCCCTGAAGGAGTCGGGCTCCGAGAAGCCCAG CGAGGACGTCCTGGTCAAGGTGTTTGAGAAGTTCGGGGAGATCCGGAATGTGGACATCCCCATGCTGGACCCCTA CCGGGAGGAGATGACGGCCGCAACTTCCACACCTTCAGTTTCGGGGGGCACTTGAACTTCGAGGCCTATGTGCA GTACCGTGAGTACATGGGCTTCATCCAGGCCATGAGCGCCCTGCGCGGGATGAAACTCATGTACAAGGGCGAGGA CGGCAAGGCCGTGGCCTGCAACATCAAGGTTTCTTTTGATTCGACCAAACACCTGAGTGATGCCTCAATTAAGAA GCGGCAGCTGGAGAGGCAGAAGCTTCAGGAACTGGAGCAGAGAAGAAGAAAAAAAGCGCAGAGAAGAAGCAAAGC GAAGCTTCGCAAGAGGGACCAGAAGCAGAGGGACCGTGAGCTGCGCCGGAATCAGAAGAAGCTGGAGAAGCTGCA GGCGGAGGAGCAGAAGCAGCTGCAGGAGAAGATCAAGCTGGAGGAGCGCAAGCTGCTGCTGCCCAGAGGAACCT GCAGTCCATCCGCCTCATCGCCGAGCTGCTCAGCAGAGCCAAGTTGTGATAAGTCCCCGGGAAGGAGCATGACAA GAGGCTGAGACATGTGGCAGCCCAGTCCTTACCTCATGGCTGTGAAGCTACGGGAACAGGAGCAGAAGGAGGAGAAG AGCGCGCGCTGGGCCTGCAGCGGAAAGAGCGGGAGCTGCGCGAGCGGCTGCTGAGCATCCTGCTGAGCAAGAAGC CGGACGACACACACACGACGAGCTGGGCGTGGCACACGCCGACCTGCTGCAGCCCGTCCTGGACATCCTGC AGGAGAGCGCGCCCACCCAGAGGCCGACGGCGCTCCCAAAAGCGTGAACGGGAGCGTGGCCGAGGAGGCCCCAT GCAAGGAGGTTCAGAGCTCCTGTCGTGTGGTCCCCGAGGATGGCTCTCCAGAGAAGAGGTGCCCGGGCGGCGTCC CCCGGTCAGAACAGGACAAGTGCAACCGGGAGCCCAGCAAGGGCCGGGGCCGGGCCACCGGAGACGGGCTTGCTG GGCCCCACAGAAGCACGCCTACAAGGATGACAGCCCCCGCCGGCGCAGCACGAGCCCGGACCACACCCGGTCCC GCCGCCACCGCCGAAGCGAGCGGTCGCGCTCCCGGTCCCCGAGCAGGCACCGCAGTACCTGGAACAGGTAAT GACGGGCACGGCCTCCCCACGGCCTGTCCGGGAAAGACCAGGACCTGCTCGAGCCTCCTGGCCGCTCCTTGGCCG GACACCATGCTTTAGAGATCCATCTTTCTCCACTCACCGCAGCGTACTTGGCACTTCAGTTTCAAACACGTAGTC TGAGGATGGCAGAGCTGCTGCATTCCCCCACACGGGGATTTCTGTGTCTGCTTGGCGACCTCCCTGCGTGCACGG GGTACAAAATGGTCTGATGCTCCTTCAAAAACATTCACTTTTTACAACGTCAAGGAATTAAGCATAAAAAAGATT ACTTGGAATGTTTCAAACAGATTTCAGGCTGCAAACTTGTTTTATAATCGTTTGCTTCTCCAAGTGAAGCTCAGA AATACCTAAAAATAGCTGTAACGTTCGCGTTAGGAAAGATGGTGTTTATTCCAGTTTGCATTTTTATGGTGAAAT AAAATCCG

875/2825 FIGURE 768

MAAATIVHDTSEAVELCPAYGLYLKPITKMTISVALPQLKQPGKSISNWEVMERLKGMVQNHQFSTLRISKSTMD FIRFEGEVENKSLVKSFLACLDGKTIKLSGFSDILKVRAAEFKIDFPTRHDWDSFFRDAKDMNETLPGERPDTIH LEGLPCKWFALKESGSEKPSEDVLVKVFEKFGEIRNVDIPMLDPYREEMTGRNFHTFSFGGHLNFEAYVQYREYM GFIQAMSALRGMKLMYKGEDGKAVACNIKVSFDSTKHLSDASIKKRQLERQKLQELEQQREEQKRREKEAEERQR AEERKQKELEELERERKREEKLRKREQKQRDRELRRNQKKLEKLQAEEQKQLQEKIKLEERKLLLAQRNLQSIRL IAELLSRAKL

876/2825 FIGURE **769**

TCTCTCGCGGCCTCAAAGCGCGGCCTGCGTCGCTTCCGGCAGTTCCAGACCGCGGGCGATGCCGCTGGGGG CGCTGGATCCAGCGGCCTGCTGAGGAACCGGGGGCCGGGCGGAGCCGGAGCCGGTCGCTGAGCGTGTC GGCGCGGGCCCGGAGCAGCTCAGAAGATAAAATAACAGTCCACTTTATAAACCGTGATGGTGAAACATTAACAAC CAAAGGAAAAGTTGGTGATTCTCTGCTAGATGTTGTGGTTGAAAATAATCTAGATATTGATGGCTTTGGTGCATG TGAGGGAACCCTGGCTTGTTCAACCTGTCACCTCATCTTTGAAGATCACATATATGAGAAGTTAGATGCAATCAC GACAAAATCTATGGACAATATGACTGTTCGAGTGCCTGAAACAGTGGCTGATGCCAGACAATCCATTGATGTGGG CAAGACCTCCTGAACTAGAACAAATAGGAATATTTTCATGGAATTTTTACCTATTTTTATAATTATTATTTCTTAA AGTGATTAAATGAGAACATGGATGAGTGGACTTCATATTATGACTAGCTTTACTATTTTAATTCACCTTGCATAA TCATATATCTCATCTGTTTACCTAGAAGATAGGTTAAGGAAATATATTATTATTCCTGTTTGATGTGGGTGAAG GCAGAGATCTAACCTGGCTTGTTTAGGGCCATACCACTAATTAGAAAATCTGTGCTAGAACCTGTGTCTTATTCC TTTCAAAATTTTGATTCGGAAGACTAAGTCTGGAGCTAGACATTATAATGCTATCAAAGAAGTTGATCTCTGTTT TGACTAAACTAGAGGAAAAATGATTGGATGAGTTTATTCTTTTCTAAGCAGAATGGTTTAACTTTGTACTCAAAG AAAAATAATGCTGATTTATAAATCTCTGCCTATAATAGAATGG

877/2825 FIGURE 770

MAAAGGARLLRAASAVLGGPAGRWLHHAGSRAGSSGLLRNRGPGGSAEASRSLSVSARARSSSEDKITVHFINRD GETLTTKGKVGDSLLDVVVENNLDIDGFGACEGTLACSTCHLIFEDHIYEKLDAITDEENDMLDLAYGLTDRSRL GCQICLTKSMDNMTVRVPETVADARQSIDVGKTS

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878/2825 FIGURE **77**1

 $\verb|CTTCTCTTTTCCCTAGACTGCAGCCAGCGGAGCCCGCAGCCAGGCCAGGAACCCAGGTCCGGAGCCTCAA| \\$ $\tt CTTCAGG{\color{red} ATC} TTGACAACATTGCTGCCGATACTGCTGCTGTCTGGCTGGGCCTTTTGTAGCCAAGACGCCTCAGA$ TGGCCTCCAAAGACTTCATATGCTCCAGATCTCCTACTTCCGCGACCCCTATCACGTGTGGTACCAGGGCAACGC GCAGGAGCCCGAGAGCTGGGCGCACGCAGAGTGGCCTGCAGTCCTACCTGCTCCAGTTCCACGGCCTCGTGCG $\tt CCTGGTGCACCAGGAGCGGACCTTGGCCTTTCCTCTGACCATCCGCTGCTTCCTGGGCTGTGAGCTGCCTCCCGA$ $\verb|CTTGTGGCAGGCAGACACCCAGGTCACCTCCGGAGTGGTCACCTTCACCCTGCAGCAGCTCAATGCCTACAACCG|\\$ CACTCGGTATGAACTGCGGGAATTCCTGGAGGACACCTGTGTGCAGTATGTGCAGAAACATATTTCCGCGGAAAA CACGAAAGGGAGCCAAACAAGCCGCTCCTACACTTCGCTGGTCCTGGGCGTCCTGGTGGGCGGTTTCATCATTGC CTGGATTGATGGAGGCTGGCAAGGGAAAGTTTCAGCTCACTGTGAAGCCCAGACTCCCCAACTGAAACACCAGAAG GTGGACAAAGTACTTGGTTTGCTAAGAACCTAAGAACGTGTATGCTTTGCTGAATTAGTCTGATAAGTGAATGTT TCACCTGAGGCGTTCAAAAGATATAACCAAATAAACAAGTCATCCACAATCAAAATACAACATTCAATACTTCCA AAATCCAAGTCATATGGCAGTGATCAATTATTAATCAATTAATAATATTAATAAATTTCTTATATTTAAGGCAAA AAAAAAAAAAAAAAAAAAAAAAAAAAAA

879/2825 **FIGURE 772**

MLTTLLPILLLSGWAFCSQDASDGLQRLHMLQISYFRDPYHVWYQGNASLGGHLTHVLEGPDTNTTIIQLQPLQE PESWARTQSGLQSYLLQFHGLVRLVHQERTLAFPLTIRCFLGCELPPEGSRAHVFFEVAVNGSSFVSFRPERALW QADTQVTSGVVTFTLQQLNAYNRTRYELREFLEDTCVQYVQKHISAENTKGSQTSRSYTSLVLGVLVGGFIIAGV AVGIFLCTGGRRC

880/2825 FIGURE 773A

ACTCCCGGCCGGGGTAGCTCTTCACTCCTCAGCGCGACGTCGTGTCGAGTTCCCAAAAAGCTCCGCAGGGGCTGT GCATTCCAGATTCTTATATTCAGCTGCCTTTTGGATATCTCTCCCAGGATGTTCTCAAGGCATACAAGAATTAAA TTCTGAATAAGTCTGCAGGTAGGATGGACAGTTATTTTAAAGCAGCTGTCAGTGACTTGGACAAACTCCTTGATG ATTTTGAACAGAACCCAGATGAACAAGATTATCTCCAAGATGTACAAAATGCATATGATTCTAACCACTGCTCAG TTTCTTCAGAGTTGGCTTCCTCACAGCGAACTTCATTGCTCCCAAAAGACCAAGAGTGCGTTAATAGTTGTGCCT ATGAAAAAATGTAACAGGACTTGATCTTCTTCTTCTGTGGATGGTGGTACTTCAGATGAAATCCAGCCGTTAT ATATGGGACGATGTAGTAAACCTATCTGTGATCTGATAAGTGACATGGGTAACTTAGTTCATGCAACCAATAGTG AAGAAGATATTAAAAAATTATTGCCAGATGATTTTAAGTCTAATGCAGATTCCTTGATTGGATTGGATTTATCTT CAGTGTCAGATACTCCCTGTGTTTCTTCAACAGACCATGATAGTGATACTGTCAGAGAACAACAGAATGATACCA GTTCTGAATTACAAAATAGAGAAATCGGAGGAATCAAAGAATTGGGTATAAAAGTAGATACAACACTTTCAGATT CCTATAATTACAGTGGAACAGAAAATTTAAAAGATAAAAAGATCTTTAATCAGTTAGAATCAATTGTTGATTTTA ACATGTCATCTGCTTTGACTCGACAAAGTTCCAAAATGTTTCATGCCAAAGACAAGCTACAACACAAGAGCCAGC CATGTGGATTACTAAAAGATGTTGGCTTAGTAAAAGAGGAAGTAGATGTGGCAGTCATAACTGCCGCAGAATGTT TAAAAGAAGAGGGCAAGACAAGTGCTTTGACCTGCAGCCTTCCGAAAAATGAAGATTTATGCTTAAATGATTCAA ATTCAAGAGATGAAAATTTCAAATTACCTGACTTTTCCTTTCAGGAAGATAAGACTGTTATAAAACAATCTGCAC AAGAAGACTCAAAAAGTTTAGACCTTAAGGATAATGATGTAATCCAAGATTCCTCTTCAGCTTTACATGTTTCCA CACGGGGTGATTTTTTACCTCAGCATGAACATAAAGATAATATACAAGATGCAGTGACTATACATGAAGAAAATAC AGAACAGTGTTGTTCTAGGTGGGGAACCATTCAAAGAGAATGATCTTTTGAAACAGGAAAAATGTAAAAGCATAC TGGATGGTGGTGACACCAGTTCTACAGTTGTAGAATCTCAAGAGGGGCTTTCTGGCACTCATGTCCCAGAGTCTT TTGATGAAGGCGCAAAAAGTGGCCCACTAATTAGTGATGCTGAACTTGATGCCTTTCTGACAGAACAGTATCTTC GCTTAGATGATGGAAACATCAATAATATATATTTCAATGCAGAAGCAGGAGCTATTGGGGAAAGTCATGGTATTA CAGTTCAACAAGGGTTACCTACCAGTAAGTCTGAGATTACAAATCAATTATCAGTCTCTGATATTAACAGTCAATCAGATGTTCCAGATACAATAGAAAGTGAACCCAGCACAGCAGATACCGTTGTTCCAATCACTTGTGCTATAGATT $\verb|CTACAGCTGATCCACAGGTTAGCTTCAACTCTAATTACATTGATATAGAAAGTAATTCTGAAGGTGGATCTAGTT|\\$ CTTGGGTTCCTGATTCAGAAGCTCCAAACTGTATGAACTGCCAAGTCAAATTTACTTTTACCAAACGGCGACACC ATTGCCGAGCATGTGGGAAAGTATTTTGTGGTGTCTGTTGTAATAGGAAGTGTAAACTGCAATATCTAGAAAAGG AAGCAAGAGTATGTGTAGTCTGCTATGAAACTATTAGTAAAGCTCAGGCATTTGAAAGGATGATGAGTCCAACTG GTTCTAATCTTAAGTCTAATCATTCTGATGAATGTACTACTGTCCAGCCTCCTCAGGAGAACCAAACATCCAGTA TACCTTCACCAGCAACTTTGCCAGTCTCAGCACTTAAACAACCAGGTGTTGAAGGACTATGTTCCAAAGAACAGA AGAGAGTATGGTTTGCAGATGGTATATTGCCCAATGGTGAAGTTGCAGATACAACAAAATTATCATCTGGAAGTA AAAGATGTTCTGAAGACTTTAGTCCTCTCTCACCTGATGTGCCTATGACAGTAAACACAGTGGATCATTCCCATT CTACTACAGTGGAAAAGCCAAACAATGAGACAGGAGATATTACAAGAAATGAGATAATTCAGAGTCCTATTTCTC AGGTTCCATCAGTGGAAAAATTGTCTATGAACACAGGAAATGAGGGGTTACCTACTTCTGGTTCATTTACACTAG ATGATGATGTTTTTGCAGAAACTGAAGAACCATCTAGTCCTACTGGTGTCTTAGTTAACAGCAATTTACCTATTG CTAGTATTTCAGATTATAGGTTACTGTGTGATATTAACAAGTATGTCTGCAATAAGATTAGTCTTCTACCTAATG ATGAGGACAGTTTGCCCCCACTTCTGGTTGCATCTGGAGAAAAGGGATCAGTGCCTGTAGTAGAAGAACATCCAT CTCATGAGCAGATCATTTTGCTTCTTGAAGGTGAAGGCTTTCATCCTGTTACATTTGTCCTAAATGCTAATCTAC ${\tt TCGTGAATGTCAAATTCATATTTTATTCCTCAGACAAATATTGGTACTTTTCAACCAATGGATTGCATGGCTTGG}$ GACAGGCAGAAATTATTATTCTATTGTTATGTTTGCCAAATGAAGATACTATTCCTAAGGACATCTTCAGACTAT

881/2825 FIGURE 773B

TTATCACCATATATAAGGATGCTCTAAAAGGAAAAATACATAGAAAAACTTGGACAATATTACCTTTACTGAGAGTT TTCTCAGTAGCAAGGATCACGGAGGATTCCTGTTTATTACACCTACTTTTCAGAAACTTGATGATCTCTCATTAC CAAGTAATCCTTTTCTTTGTGGAATTCTTATCCAGAAGCTTGAGATTCCCTGGGCAAAGGTTTTTCCTATGCGTT TAATGTTGAGATTGGGTGCAGAATATAAAGCATATCCTGCTCCTCTAACAAGCATCAGAGGCCGAAAACCTCTTT TTGGAGAAATAGGACACACTATTATGAACTTACTTGTTGACCTTCGAAATTACCAGTATACCTTGCATAATATAG ATCAACTGTTGATTCATATGGAAATGGGAAAAAGCTGCATAAAAATACCACGGAAAAAGTACAGTGATGTAATGA AAGTACTAAATTCTTCCAATGAGCATGTCATTAGCATTGGAGCAAGTTTCAGTACAGAAGCAGATTCTCATCTAG TCTGTATACAGAATGATGGAATTTATGAAACACAGGCCAACAGTGCCACTGGCCATCCTAGAAAAGTGACAGGTG CAAGTTTTGTGGTATTCAATGGAGCTCTAAAAACATCTTCAGGATTTCTTGCTAAGTCCAGCATAGTTGAAGATG GCTTAATGGTACAAATAACTCCAGAGACCATGAATGGCTTGCGGCTAGCTTTACGAGAACAGAAAGACTTTAAAA TTACATGTGGGAAAGTTGATGCAGTAGACCTGAGAGAATACGTGGATATCTGCTGGGTAGATGCTGAAGAAAAAG GAAACAAAGGAGTTATCAGTTCAGTGGATGGAATATCATTACAAGGATTTCCAAGTGAAAAAATAAAACTGGAAG CAGATTTTGAAACCGATGAGAAGATTGTAAAATGTACCGAGGTGTTCTACTTTCTAAAGGACCAGGATTTATCTA TTTTATCAACTTCTTATCAGTTTGCAAAAGAAATAGCCATGGCTTGTAGTGCTGCGCTGTGCCCTCACCTGAAAA CTCTAAAAAGTAATGGGATGAATAAAATTGGACTCAGAGTTTCCATTGACACTGATATGGTTGAATTTCAGGCAG GATCTGAAGGCCAACTTCTGCCTCAGCATTATCTAAATGATCTTGATAGTGCTCTGATACCTGTGATCCATGGTG GGACCTCCAACTCTAGTTTACCATTAGAAATAGAATTAGTGTTTTTCATTATAGAACATCTTTTT**TAG**TGAAAGA AAAAGTATAAATATGTCTGATTTTTGAAACACATAAGCTTTGCTCTTTTAGGCAGGAATGATCTTTTCAAATCATT AAGACAAGGATTTCTTTTAAGAAATTTATAGCATTTACTGTGTTATTTAAATGCTAAGCCAAAGTATCTGCACTT AGGTATACCTCTTTATGCCAATAATGATTTTAATGAAGGCTCTTTTCAGATGTAACCTTATGAAGGAAATATCTG CTTTGTGTATATGCCAGTTAGAATACTGGTTTCTAAAGTCTGTCAAATTGTATTTCAGTGGCACAAAAACCAGTT ATTATATATCCACATATATAGTTTTCCCTGATTAAATGGATATTAAAATAATTGCGGGTGCTTCAGGACTTTTTG CTTCTATATTTAAGTATATTGTTTTTATAGCAAGAACATATTCTGAATGTTTTATAAATCTTTAATAATTTATAT GTAGGTAATATTTTTGTATCACAATGCATTATTTTTTTTCCTCCTTTCCAAACTATACCACTGTATTTACC ACTTCTAAGAGTGACTGACGGGCCAGATGACCCTTGAAGTAGTCATTATGTAGCAATAAATGAAGCCTGAAA CAGGTTTTTTTACTTCCACTTTAATCCTTAGAAATTTCTTGGCAACTTCGCATATTTTCATTGACACTGGTGTAT AAGTATAAATTTAAATGAACTAATTACTTTTGCATATTTTAAATTCTTTATATGGTAGTTATTTTTTATAACAGG ATATTAACATAAGTTAAATCCTATGTATTTGAAATTGTTACAGAGCTTTCCTCTTTACTTCAAACAGCAAAAAAG TGGGGGGCATATTGTAGTCCTGTCATTTAAGTTATGTAAAAAAATTTAATCATTATTTTGATGCTTTAAACATTCT ${\tt CATGTGTAATATGTTTTTGTATCAAAAACACTCATATATTTCAAGAAAAAGAAATTATGTTAAATAGCCCTGT}$ TTTAAGAAAATATTTATGAAGCATCTCAACTTGAAGATCAAGTCAAAGTTATAACTCAGGATCTGAGGTCTCAA GCTAGGAGAGACTGAGAATTTTAATCAGTTTGGGCATATAGTTTGGACTGAATCACATCTGTAGTACTTAGCCAA AGACAATTTGGAGGAGAATATCAGCCTTCTGGAAGTAGCTACTTCCTGAACAATGTAAAGTGTCGCAGATATTCA ATAAAATGGCAACCTGTTATAATTTGTGAAATTTATTGAAATGGTGTAAGATGAAAACAATTGCATATCAAACCC AGAGGGCTGTTCCATGATGGGACAGCTTTGGATTTGTTTTCATAAAATCTCTACATTCAATAAAAATTGGAATTA TGTGCCTGAAGTTTGGAGGCACATTTTGAAGT

882/2825 FIGURE **774**

MDSYFKAAVSDLDKLLDDFEQNPDEQDYLQDVQNAYDSNHCSVSSELASSQRTSLLPKDQECVNSCASSETSYGT NESSLNEKTLKGLTSIQNEKNVTGLDLLSSVDGGTSDEIQPLYMGRCSKPICDLISDMGNLVHATNSEEDIKKLL PDDFKSNADSLIGLDLSSVSDTPCVSSTDHDSDTVREQQNDTSSELQNREIGGIKELGIKVDTTLSDSYNYSGTE NLKDKKIFNQLESIVDFNMSSALTRQSSKMFHAKDKLQHKSQPCGLLKDVGLVKEEVDVAVITAAECLKEEGKTS ALTCSLPKNEDLCLNDSNSRDENFKLPDFSFQEDKTVIKQSAQEDSKSLDLKDNDVIQDSSSALHVSSKDVPSSL SCLPASGSMCGSLIESKARGDFLPQHEHKDNIQDAVTIHEEIQNSVVLGGEPFKENDLLKQEKCKSILLOSLIEG MEDRKIDPDQTVIRAESLDGGDTSSTVVESQEGLSGTHVPESSDCCEGFINTFSSNDMDGODLDYFNIDEGAKSG PLISDAELDAFLTEQYLQTTNIKSFEENVNDSKSQMNQIDMKGLDDGNINNIYFNAEAGAIGESHGINIICETVD KQNTIENGLSLGEKSTIPVQQGLPTSKSEITNQLSVSDINSQSVGGARPKQLFSLPSRTRSSKDLNKPDVPDTIE SEPSTADTVVPITCAIDSTADPQVSFNSNYIDIESNSEGGSSFVTANEDSVPENTCKEGLVLGQKQPTWVPDSEA PNCMNCQVKFTFTKRRHHCRACGKVFCGVCCNRKCKLQYLEKEARVCVVCYETISKAQAFERMMSPTGSNLKSNH SDECTTVQPPQENQTSSIPSPATLPVSALKQPGVEGLCSKEQKRVWFADGILPNGEVADTTKLSSGSKRCSEDFS PLSPDVPMTVNTVDHSHSTTVEKPNNETGDITRNEIIQSPISQVPSVEKLSMNTGNEGLPTSGSFTLDDDVFAET EEPSSPTGVLVNSNLPIASISDYRLLCDINKYVCNKISLLPNDEDSLPPLLVASGEKGSVPVVEEHPSHEQIILL LEGEGFHPVTFVLNANLLVNVKFIFYSSDKYWYFSTNGLHGLGOAEIIILLLCLPNEDTIPKDIFRLFITIYKDA LKGKYIENLDNITFTESFLSSKDHGGFLFITPTFQKLDDLSLPSNPFLCGILIQKLEIPWAKVFPMRLMLRLGAE YKAYPAPLTSIRGRKPLFGEIGHTIMNLLVDLRNYQYTLHNIDQLLIHMEMGKSCIKIPRKKYSDVMKVLNSSNE HVISIGASFSTEADSHLVCIQNDGIYETQANSATGHPRKVTGASFVVFNGALKTSSGFLAKSSIVEDGLMVQITP ETMNGLRLALREQKDFKITCGKVDAVDLREYVDICWVDAEEKGNKGVISSVDGISLQGFPSEKIKLEADFETDEK IVKCTEVFYFLKDQDLSILSTSYQFAKEIAMACSAALCPHLKTLKSNGMNKIGLRVSIDTDMVEFQAGSEGQLLP QHYLNDLDSALIPVIHGGTSNSSLPLEIELVFFIIEHLF

883/2825 FIGURE 775

GGCACGAGGGCGATGCGGGCGCCCCCGGCGGGCGGCCCCGGCGGCACCATCAGCCCTCTGCTCCGCCGCCTGCT GCTCGCCGCACTCCTGCAGCTGGCCCCCGCCCAGGCCCTGTCTCCCAGCCTGATGCCCCTGGCCACCAGAGGAA AGTGGTGTCATGGATAGATGTGTATACTCGCGCTACCTGCCAGCCCCGGGAGGTGGTGGTGCCCTTGACTGTGGA GCTCATGGGCACCGTGGCCAAACAGCTGGTGCCCAGCTGCGTGACTGTGCAGCGCTGTGGTGGCTGCCCTGA CGATGGCCTGGAGTGTGCCCCACTGGGCAGCACCAAGTCCGGATGCAGATCCTCATGATCCGGTACCCGAGCAG GCCAGACAGGGCTGCCACTCCCCACCACCGTCCCGGTTCTGTTCCGGGCTGGGACTCTGCCCCCGGAGC ACCCTCCCCAGCTGACATCACCCATCCCACTCCAGCCCCAGGCCCCTCTGCCCACGCTGCACCCAGCCACCAG $\tt CGCCCTGACCCCGGGACCTGCCGCTGCCGACGCCGCAGCTTCCTCCGTTGCCAAGGGCGGGGCT{\color{red}{\bf TAG}} AG$ CTCAACCCAGACACCTGCAGGTGCCGGAAGCTGCGAAGGTGACACATGGCTTTTCAGACTCAGCAGGGTGACTTG CCTCAGAGGCTATATCCCAGTGGGGGAACAAAGAGGGGCCTGGTAAAAAACAGCCAAGCCCCCAAGACCTCAGCC CAGGCAGAAGCTGCTCTAGGACCTGGGCCTCTCAGAGGGCTCTTCTGCCATCCCTTGTCTCCCTGAGGCCATCAT CAAACAGGACAGAGTTGGAAGAGGAGACTGGGAGGCAGCAAGAGGGGTCACATACCAGCTCAGGGGAGAATGGAG ${\tt TACTGTCTCAGTTTCTAACCACTCTGTGCAAGTAAGCATCTTACAACTGGCTCTTCCTCCCCTCACTAAGAAGAC}$ CCAAACCTCTGCATAATGGGATTTGGGCTTTGGTACAAGAACTGTGACCCCCAACCCTGATAAAAGAGATGGAAG

884/2825 FIGURE 776

 ${\tt MSPLLRRLLLAALLQLAPAQAPVSQPDAPGHQRKVVSWIDVYTRATCQPREVVVPLTVELMGTVAKQLVPSCVTV} \\ {\tt QRCGGCCPDDGLECVPTGQHQVRMQILMIRYPSSQLGEMSLEEHSQCECRPKKKDSAVKPDRAATPHHRPQPRSV} \\ {\tt PGWDSAPGAPSPADITHPTPAPGPSAHAAPSTTSALTPGPAAAAADAAASSVAKGGA} \\ {\tt CRCGGCPDDGLECVPTGQHQVRMQILMIRYPSSQLGEMSLEEHSQCECRPKKKDSAVKPDRAATPHHRPQPRSV} \\ {\tt PGWDSAPGAPSPADITHPTPAPGPSAHAAPSTTSALTPGPAAAAADAAASSVAKGGA} \\ {\tt CRCGGCPDDGLECVPTGQHQVRMQILMIRYPSSQLGEMSLEEHSQCECRPKKKDSAVKPDRAATPHHRPQPRSV} \\ {\tt CRCGGCPDGGAPSPADITHPTPAPGPSAHAAPSTTSALTPGPAAAAADAAASSVAKGGA} \\ {\tt CRCGGCPDGGAPSPADITHPTPAPGPSAHAAPSTTSALTPGPAAAAADAAASSVAKGGA} \\ {\tt CRCGGCPDGGAPSPADITHPTPAPGPSAHAAPSTTSALTPGPAAAAADAAASSVAKGGA} \\ {\tt CRCGGCPDGGAPSPADITHPTPAPGPSAHAAPSTTSALTPGPAAAAADAAASSVAKGGA} \\ {\tt CRCGGCPDGGAPSPADITHPTPAPGPSAHAAPSTTSALTPGPAAAAAADAAASSVAKGGA} \\ {\tt CRCGGCPDGGAPSTTAATGAPTAATGAPTA$

885/2825 FIGURE 777

GCAGCCCACTCTTCCCGAGAGGCCCCGACATCTCCCGGATAATCTGGAGCTTTCAAAATCTTTGCCAAAGCAACC AGGAATTTTCTTAGAGAAGTTGATGCTGATGGTGACCTGATTGCAGTATCAAATCTGAATGACTCTGATAAGTTA CAGCTTCTAAGTCTGGTGACAAAAAAGAAGAGATTCTGGTGCTGGCAGAGACCCAAGTACCAGTTTTTATCCCTC ACCCTTGGCGATGTACTCATAGAAGACCAATTTCCGAGTCCAGTGGTCGTGGAGTCGGACTTTGTGAAATACGAG GGCAAGTTTGCAAACCACGTGAGTGGAACCCTGGAGACTGCACTGGGGAAGGTCAAGCTGAACCTGGGGGGCAGC AGCCGCGTAGAGAGCCAGTCTTCATTTGGAACCCTGAGGAAGCAGGAGGTGGATTTGCAGCAGCTCATCAGAGAC GTTTTGACACAGAAGATCACGACGATGCAGAAGTGTGTGATCTCTGAGCACATGCAGGTCGAGGAGAAGTGTGGT GGCATCGTGGGCATCCAGACCAAGACGGTGCAGGTGTCAGCGACGGAGGATGGGAATGTCACCAAGGACTCCAAC GTGGTGCTGGAGATCCCAGCTGCCACCACTTGCCTACGGTGTCATTGAGTTATACGTGAAACTGGACGGCCAG TTCGAGTTCTGCCTTCTCCGAGGGAAGCAAGGTGGCTTCGAGAACAAGAAGAAATTGACTCTGTCTACCTGGAC CCCCTGGTCTTTCGAGAGTTTGCATTCATAGACATGCCAGATGCTGCGCATGGGATATCTTCCCAGGATGGACCA CAGACAGCTTTGAGTGACATCTTCCAGGCGGTCCTATTTGATGATGAACTACTCATGGTCCTGGAACCAGTGTGC GATGACCTGGTCAGCGGCCTCTCGCCCACAGTGGCGGTGCTGGGGGAGCTGAAGCCCCGGCAGCAGCACCATT CTGTTTATGACAGCCTACTTCTTGGTCAGTGCCCTCGCAGAAATGCCAGATAGCGCGGCAGCTCTGCTGGGCACT TGCTGCAAACTCCAGATCATTCCCACACTGTGCCACTTGCTTCGTGCTCTGATGATGGAGTATCTGATCTT GAAGACCCAACCTTGACTCCCCTGAAAGATACAGAAAGGTTTGGGATTGTGCAGCGCTTGTTTGCCTCAGCTGAC ATTAGTCTGGAGAGACTGAAGTCATCTGTGAAAGCTGTCATTCTGAAGGACTCTAAAGTCTTCCCACTGCTTCTT TGTATAACCCTGAATGGACTCTGTGCTTTAGGCAGAGAACATTCA**TGA**TGTCATATGTGAACTAGAAGTACGTGT TACTGGCCAAGGCTATTTTTCAGAACTGTTAAAGGTCATATGCACGTTAAAAGTTGACCAATGAAATGAATTTAC TTTTCCTTATCAGAGGCCCTCCTGTGCTGGTGACCCAGCATCTGAGTTAGGTTCCAGCATGTAAAGAGCTGGGAG GGCGGAGAATTCTTAGCATACATTCAGACGTTTTTTCTGCACAATAATAAGTCCATCTGTCACTTGCATTCCACT TTTTGTTACATAGAAAGAGTCTGACCCTTTAATCCAAAAGGTCTTTTTACATTGTGAATGCTGTGGGAAGGCAAT TTCTCTGCACAAGAGGCTACGTTTTGGAAGTGATGTTATTTGATGACTGAAAATGAACTGTAAATGCTC CTAGAGTATATTCCTCTGCTGAACAAATTAAACTTCAAAAAAATCTAACAGTAACACCCCTGCTTGGGACCC TAGCTATATGCATTTTATGTGACCTTGCCATGCTTCAGTGAACATACTAATTCTATGTCTAGCACATGTTGATTT

886/2825 FIGURE 778

MFAKATRNFLREVDADGDLIAVSNLNDSDKLQLLSLVTKKKRFWCWQRPKYQFLSLTLGDVLIEDQFPSPVVVES
DFVKYEGKFANHVSGTLETALGKVKLNLGGSSRVESQSSFGTLRKQEVDLQQLIRDSAERTINLRNPVLQQVLEG
RNEVLCVLTQKITTMQKCVISEHMQVEEKCGGIVGIQTKTVQVSATEDGNVTKDSNVVLEIPAATTIAYGVIELY
VKLDGQFEFCLLRGKQGGFENKKRIDSVYLDPLVFREFAFIDMPDAAHGISSQDGPLSVLKQATLLLERNFHPFA
ELPEPQQTALSDIFQAVLFDDELLMVLEPVCDDLVSGLSPTVAVLGELKPRQQQDLVAFLQLVGCSLQGGCPGPE
DAGSKQLFMTAYFLVSALAEMPDSAAALLGTCCKLQIIPTLCHLLRALSDDGVSDLEDPTLTPLKDTERFGIVQR
LFASADISLERLKSSVKAVILKDSKVFPLLLCITLNGLCALGREHS

887/2825 FIGURE 779

888/2825 FIGURE 780

MSLLLLVVSALHILILILLFVATLDKSWWTLPGKESLNLWYDCTWNNDTKTWACSNVSENGWLKAVQVLMVLSLI LCCLSFILFMFQLYTMRRGGLFYATGLCQLCTSVAVFTGALIYAIHAEEILEKHPRGGSFGYCFALAWVAFPLAL VSGIIYIHLRKRE

889/2825 FIGURE **781**

CCGAGAGGCCCCTGCCCAACCCGGAGGGACTGGATAGCGACTTCCTTGCCGTGCTAAGTGACTACCCGTCTCCTG ACATCAGCCCCCGATATTCCGCCGAGGGGAGAAACTGCGTGTGATTTCTGATGAAGGGGGGCTGGTGGAAAGCTA AGGGCCTGGGCAGAGACAAGGCCGAGGAGCTGCTGCAGCTGCCAGACACAAAGGTCGGCTCCTTCATGATCAGAG TGCCGAACAACTGGTACTACATTTCCCCGAGGCTCACCTTCCAGTGCCTGGAGGACCTGGTGAACCACTATTCTG CCTCCAGCTCACCTGTCACCTTGCGTCAGAAGACTGTGGACTGGAGGAGGAGGAGGACCCCG AGGGAACAGAGAACCCGCTTGGGGTAGACGAGTCCCTTTTCAGCTATGGCCTTCGAGAGAGCATTGCCTCTTACC TGTCCCTGACCAGTGAGGACAACACCTCCTTTGATCGAAAGAAGAAAGCATCTCCCTGATGTATGGTGGCAGCA ${\tt AGAGAAAGAGCTCATTCTCATCACCACCTTACTTTGAGGAC} {\color{red} {\bf TAG} {\tt CCAAGAACAGACACAATGGTTCATGCCC} }$ AAAAGGAACAGAAGTTCCAACTATTGCCTGGGATCTTGCGAAAAGCGAGGTTCCCTGATCCCTGGGAGCCTCACG TATTTTAGAAGCCAAGAGAAGCCACATGGAGACTCAAATTCGCATCTTCTCTATCCACATCATGACCAAAGGAAC CCCTCCCTGGTGTCTGATCAGGGCTGTGGCATCACAAAACATTGGATCATGACATGTCGGGCGATGCTTGGAAAA GCCCAGCATGTATGTATGCACACATTGTGTGTGTGGGAAGGACAAAGCCACTCTCACAAGAAAGGGCACCAGGAC CTGAGAGGGAGAGCAAAAGCTGTGTCCTGGCAGAAGGTCTGGGTTTGCAGATGGGTGCCCTGAATGGAACT ACTITAACTAATCCATAGGGACTTCTGGTATGCTTTCCTCTCTTTTTAAAGGAACTTCGTGACACTAAACATTAG $\verb|CCCAAAGGACTTCTTAGCCTTCAATTGGGAGATACCTTTGGTCTGCTCCTGCACCAAAGCCATATGGGTGGAAGT|\\$ GCGAGACTAGAACTGCACCAGCTTCCCTGATGTCTGCAGCCATGGCTTTGCAGCGCAAACAGAACTTCTCTGGGA TGCTGGGATTCTTGCCTGTATGAATGCATCAAGTATTCATTTATTGCCCGAATAGGCATTGCATTAAGTCCTCTG TTAGGTGTCAGGCAAGCCAAAAAAAAAAAAAAAGATGCTAAGTCCTAACCCCCAACAGAAGTGTTCACAGTGTAGA CGGGAAAAAATGTATAAACAAATGTGTAAAAAGAGAAATCAGCTCATGGCTTAGGATGGAATTAGAGACAGGTGA TGGATTTTTTAGAAAAGAATGGATACAGTCTAAAGAATTAACTCACCCGGTACTTTATTCTAAGAAGGGTCTGG CATCCATATGAGGAAAAATGCTCAGCTCCAGGAAAGATGGGGAGTCCAAGTGGATTAATGATGTCATGCATAATT TTAAGAGACAAGGGAGAAAACACAATGTATAGCCAGAGAAGGAGAAGCTCCCATCCAAATCCTACTAGGAAGAGA GTGGGCTGCAGATGAATCTGTGACTCATGTTTCCCTGTTTCAAAGGGATCCTGGGGAAGGAGGGGAACATGCTTG CAGTATCTCTCCCTGTCTGTCTGCTCACATAAGCATTCCGTCCATCTAAGCTCATCGTGCTACTGGTATGTGTAT GTGCAGTTACACAGTTTCCTGTATCATAAATCCTAGTGTGTTTATACAAGGAGACATCTGTGGTTTCCCCAACCG ATAAGGCAGGGTGACCCCCTGAACTCCAGATGTTTGCACAGTATCTTATGTGTTGTTTTCCGTTGTGACGAATGT GATTGGAACATTTGGGGAGCACCCAGAGGGATTTTTCAGTGGGAAGCATTACACTTTGCTAAATCATGTATTTAT TCCTGATTAAAACAAACCTAATAAATATTTAACCCTTGGC

890/2825 FIGURE 782

MGNSMKSTPAPAERPLPNPEGLDSDFLAVLSDYPSPDISPPIFRRGEKLRVISDEGGWWKAISLSTGRESYIPGI CVARVYHGWLFEGLGRDKAEELLQLPDTKVGSFMIRESETKKGFYSLSVRHRQVKHYRIFRLPNNWYYISPRLTF QCLEDLVNHYSEVADGLCCVLTTPCLTQSTAAPAVRASSSPVTLRQKTVDWRRVSRLQEDPEGTENPLGVDESLF SYGLRESIASYLSLTSEDNTSFDRKKKSISLMYGGSKRKSSFFSSPPYFED

891/2825 FIGURE 783

GGCCAGGAACGCACGCTGCCTGGCCGTATCGCCGCCCCACCGACGCCGCCGCCGCGGGACTAGAAGTGAGCCGCCC GGGTCCCAAACGCCAGCCAGCCAGTCAGTGGGTCCCGCAGTCGCCCGCAACCGGGGCGAATCATGGCGGCCGCCA AGGTGGCTTTAACCAAGAGAGCAGATCCAGCTGAGCTTAGAACAATATTTTTGAAGTATGCAAGCATTGAGAAAA CAAAGACTGTGGAACTTTTAAGTGGAGTGGTGGATCAGACCAAAGATGGATTAATATCTTTTCAAGAATTTGTTG CCTTTGAATCTGTCCTGTGTGCCCCTGATGCTTTGTTTATGGTAGCCTTTCAGCTGTTTGACAAAGCTGGCAAAG GAGAAGTAACTTTTGAGGATGTTAAGCAAGTTTTTTGGACAGACCACAATTCATCAACATATTCCATTTAACTGGG TATTGGAAATACAACTGGAGCACGCAAAGCAAGCCTTTGTGCAACGGGACAATGCTAGGACTGGGAGAGTCACAG $\verb|CCATCGACTTCCGAGACATCATGGTCACCATCCGCCCCCATGTCTTGACTCCTTTTGTAGAAGAATGTCTAGTAG|\\$ CTGCTGCTGGAGGTACCACATCCCATCAAGTTAGTTTCTCCTATTTTAATGGATTTAATTCGCTCCTTAACAACA TGGAACTCATTAGAAAGATCTATAGCACTCTGGCTGGCACCAGGAAAGATGTTGAAGTGACTAAGGAGGAGTTTG AGCCAAGGGGACGTATGACCTTAGCAGACATTGAACGGATTGCTCCTCTGGAAGAGGGAACTCTGCCCTTTAACT TGGCTGAGGCCCAGAGGCCAGAAGGCCTCAGGTGATTCAGCTCGACCAGTTCTTCTACAAGTTGCAGAGTCGGCCT ACAGGTTTGGTCTGGGTTCTGTTGCTGGAGCTGTTGGAGCCACTGCTGTATCCTATCGATCTTGTAAAAACTC GAATGCAGAACCAACGATCAACTGGCTCTTTTGTGGGAGAACTCATGTATAAAAACAGCTTTGACTGTTTTAAGA AAGTGCTACGCTATGAAGGCTTCTTTGGACTGTATAGAGGTCTGTTGCCACAGTTATTGGGAGTTGCCCCAGAGA AGGCCATAAAACTTACAGTGAACGATTTTGTGAGGGATAAATTTATGCACAAAGATGGTTCGGTCCCACTTGCAG CAGAAATTCTTGCTGGAGGCTGCGCTGGAGGCTCCCAGGTGATTTTCACAAATCCTTTAGAAATCGTCAAGATCC TTGGGATCTACAAGGGTGCCAAAGCATGCTTTCTGCGGGACATTCCTTTCTCGGCCATCTACTTTCCGTGCTATG CTCATGTGAAGGCTTCCTTTGCAAATGAAGATGGGCAGGTTAGCCCAGGAAGCCTGCTCTTAGCTGGTGCCATAG CTGGTATGCCTGCAGCATCTTTAGTGACCCCTGCTGATGTTATCAAGACGAGATTACAGGTGGCTGCCCGGGCTG GCCAAACCACTTACAGCGGAGTGATAGACTGCTTTAGAAAGATACTGCGTGAAGAAGGACCAAAAGCTCTGTGGA AGGGAGCTGGTGCTCGTGTATTTCGATCCTCACCCCAGTTTGGTGTAACTTTGCTGACTTACGAATTGCTACAGC GATGGTTCTACATTGATTTTGGAGGAGTAAAACCCATGGGATCAGAGCCAGTTCCTAAATCCAGGATCAACCTGC CTGCCCGAATCCTGATCACGTTGGGGGCTACAAACTGGCAGTTGCTACATTTGCAGGGATTGAAAACAAATTTG GACTTTACCTACCTCTCTCAAGCCATCAGTATCTACCTCAAAGGCTATTGGTGGAGGCCCA**TAG**GAAGATCAGC $\verb|CCTGGGATAGTGCTGTCTTTTTGTGGGTACTGCAGTAAAGAACATCCCTCCTGGGAATGAAGCAATGCTTCATCC| \\$ $\tt CTTTTACGTCCATCTCTTGTTTAAATTCAAGTCCAGGCTTTTTTATCATGTGAAATCATTTCTTGTGGTGTT$ AAAATTGATGGGATTTGGTCAACACTAACATGATTTGGGGAAAGGAGCAAGTCAGAATAGAAATTAGTACTCCCC TCCTTGAACTAGGATTGTAGTCCCAAAGAGGCTACTGTAAGGCAATCATGGTGCTCAGAGCAGTGTTTCGTGTGT AAATATATTCAGCCTGGCTTCAAATTGTAAGCATGCACAAATTCTGTCTCTGGATTATATTATGAAGCTTTTATG TGAAACATGTTTCTTTGTAATGAAAACCACATTGGAGATGTTTAGTAATCATATTGTTACTGGTACCAAGACTAC $\verb|CCCTCTCCTGGAAGTGCTAATTTTGAAAAACTGCTCATATGATGTACATGTACTGATTACTGCCTATTTTAATAA| \\$ TGCTTTTAAAATCCACTTTATCTTGAATAATGTAAGACAAATATGTTCTGACATAAGTATTTAATTCATGTTGCC

892/2825 FIGURE 784

MAAAKVALTKRADPAELRTIFLKYASIEKNGEFFMSPNDFVTRYLNIFGESQPNPKTVELLSGVVDQTKDGLISF QEFVAFESVLCAPDALFMVAFQLFDKAGKGEVTFEDVKQVFGQTTIHQHIPFNWDSEFVQLHFGKERKRHLTYAE FTQFLLEIQLEHAKQAFVQRDNARTGRVTAIDFRDIMVTIRPHVLTPFVEECLVAAAGGTTSHQVSFSYFNGFNS LLNNMELIRKIYSTLAGTRKDVEVTKEEFVLAAQKFGQVTPMEVDILFQLADLYEPRGRMTLADIERIAPLEEGT LPFNLAEAQRQKASGDSARPVLLQVAESAYRFGLGSVAGAVGATAVYPIDLVKTRMQNQRSTGSFVGELMYKNSF DCFKKVLRYEGFFGLYRGLLPQLLGVAPEKAIKLTVNDFVRDKFMHKDGSVPLAAEILAGGCAGGSQVIFTNPLE IVKIRLQVAGEITTGPRVSALSVVRDLGFFGIYKGAKACFLRDIPFSAIYFPCYAHVKASFANEDGQVSPGSLLL AGAIAGMPAASLVTPADVIKTRLQVAARAGQTTYSGVIDCFRKILREEGPKALWKGAGARVFRSSPQFGVTLLTY ELLQRWFYIDFGGVKPMGSEPVPKSRINLPAPNPDHVGGYKLAVATFAGIENKFGLYLPLFKPSVSTSKAIGGGP

893/2825 FIGURE **785A**

GGACTGCGAAAGGAGCAGGGTTGCGGAGCTAGGGCTCCAGCCTGCGGCCGCGCATTCTTGCGTCTGGCCAGCCGC GAGCTCTAAGGGTCGGCCCCGGGTCCGCCCCCGCGGGCTCCCTGCCAGGCTCTCGCGGGCGCGCTCGGGGTGG GCCGGAGCCGCGCGCGCGCGCTGTAATCGGACACCAAGAGCGCTCGCCCCCGGCCTCCGGCCACTTTCCATTC ACTCCGAGGTGCTTGATTGAGCGACGCGGAGAAGAGCTCCGGGTGCCGCGCACTGCAGCGCTGAGATTCCTTTA CAAAGAAACTCAGAGGACCGGGAAGAAAGAATTTCACCTTTGCGACGTGCTAGAAAATAAGGTCGTCTGGGAAAA $\texttt{TATTTTCCCTTGGATATTAACTTGCATATCTGAAGAA} \underline{\textbf{ATG}} \texttt{GCATTCCGGACAATTTGCGTGTTGGATATT}$ AACCAAGACCTCTGAATACTTCAGCCTTTCCCACCATCCTTTAGACTACAGGATTTTATTAATGGATGAAGATCA GGACCGGATATATGTGGGAAGCAAAGATCACATTCTTTCCCTGAATATTAACAATATAAGTCAAGAAGCTTTGAG GAACTTTGTCCGTGTAATTCAGACTTTCAATCGCACACTTTGTATGTCTGTGGGAGTGGCGCTTTCAGTCCTGT CTGTACTTACTTGAACAGAGGGAGGAGATCAGAGGACCAAGTTTTCATGACTCCAAGTGTGAATCTGGAAA AGGACGCTGCTCTTTCAACCCCAACGTGAACACGGTGTCTGTTATGATCAATGAGGAGCTTTTCTCTGGAATGTA TATAGATTTCATGGGGACAGATGCTGCTATTTTTCGAAGTTTAACCAAGAGGAATGCGGTCAGAACTGATCAACA TAATTCCAAATGGCTAAGTGAACCTATGTTTGTAGATGCACATGTCATCCCAGATGGTACTGATCCAAATGATGC TCGAATATGTCCTAATGACACTGGTGGACTGCGTAGCCTTGTCAACAAGTGGACCACTTTCTTAAAGGCGAGGCT GGTGTGCTCGGTAACAGATGAAGACGGCCCAGAAACACACTTTGATGAATTAGAGGATGTGTTTCTGCTGGAAAC TGATAACCCGAGGACAACACTAGTGTATGGCATTTTTACAACATCAAGCTCAGTTTTCAAAGGATCAGCCGTGTG TGTGTATCATTTATCTGATATACAGACTGTGTTTAATGGGCCTTTTGCCCACAAAGAAGGGCCCAATCATCAGCT GATTTCCTATCAGGGCAGAATTCCATATCCTCGCCCTGGAACTTGTCCAGGAGGAGCATTTACACCCAATATGCG AACCACCAAGGAGTTCCCAGATGATGTTGTCACTTTTATTCGGAACCATCCTCTCATGTACAATTCCATCTACCC AATCCACAAAAGGCCTTTGATTGTTCGTATTGGCACTGACTACAAGTACACAAAGATAGCTGTGGATCGAGTGAA ${\tt CGCTGCTGATGGGAGATACCATGTCCTGTTTCTCGGAACAGATCGGGGTACTGTGCAAAAAGTGGTTGTTCTTCC}$ TACTAACAACTCTGTCAGTGGCGAGCTCATTCTGGAGGAGCTGGAAGTCTTTAAGAATCATGCTCCTATAACAAC AATGAAAATTTCATCTAAAAAGCAACAGTTGTATGTGAGTTCCAATGAAGGGGTTTCCCAAGTATCTCTGCACCG $\tt CTGCCACATCTATGGTACAGCCTGTGCTGACTGCTGCCTGGCGGGACCCTTATTGCGCCTGGGATGGCCATTC$ CTGTTCCAGATTCTACCCAACTGGGAAACGGAGGAGCCGAAGACAAGATGTGAGACATGGAAACCCACTGACTCA TTTTCTGGAGTGTGCCCCCAAGTCTCCGCAGGCATCTATCAAGTGGCTGTTACAGAAAGACAAAGACAGGAGGAA AGAGGTTAAGCTGAATGAACGAATAATAGCCACTTCACAGGGACTCCTGATCCGCTCTGTTCAGGGTTCTGACCA AGGACTTTATCACTGCATTGCTACAGAAAATAGTTTCAAGCAGACCATAGCCAAGATCAACTTCAAAGTTTTAGA TTCAGAAATGGTGGCTGTTGTGACGGACAAATGGTCCCCGTGGACCTGGGCCCAGCTCTGTGAGGGCTTTACCCTT CCACCCGAAGGACATCATGGGGGCATTCAGCCACTCAGAAATGCAGATGATTAACCAATACTGCAAAGACACTCG GCAGCAACATCAGCAGGAGATGAATCACAGAAAATGAGAGGGGGACTATGGCAAGTTAAAGGCCCTCATCAATAG ${\tt TCGGAAAAGTAGAAACAGGAGGAATCAGTTGCCAGAGTCA}{\tt TAA}{\tt TATTTTCTTATGTGGGTCTTATGCTTCCATTA}$ ACAAATGCTCTGTCTTCAATGATCAAATTTTTGAGCAAAGAAACTTGTGCTTTACCAAGGGGAATTACTGAAAAAG TTTCTTCAAAATCGATATGGTGGCTGCAAAACTAGAATTGTTGCATCCCTCAATTGAATGAGGGCCATATCCCTG TGGTATTCCTTTCCTGCTTTGGGGCTTTAGAATTCTAATTGTCAGTGATTTTGTATATGAAAACAAGTTCCAAAT CCACAGCTTTTACGTAGTAAAAGTCATAAATGCATATGACAGAATGGCTATCAAAAGAAATAGAAAAGGAAGACG GCATTTAAAGTTGTATAAAAACACGAGTTATTCATAAAGAGAAAATGATGAGTTTTTATGGTTCCAATGAAATAT CTTCCCCTTTTTTTAAGATTGTAAAAATAATCAGTTACTGGTATCTGTCACTGACCTTTGTTTCCTTATTCAGGA AGATAAAAATCAGTAACCTACCCCATGAAGATATTTGGTGGGAGTTATATCAGTGAAGCAGTTTGGTTTATATTC

894/2825 FIGURE 785B

TAATCTTGCACTACTCAGTTATTACTGTTTGTTCTCTTATTCCCTAGTCTGTGTGGCAAATTAAACAATATAAGA AATTTTTTAAGAGGAATTTATAACAAATATGGCTAGTTTTCAACTTCAGTACTCAAATTCAATGATTCTTCCTTT TATTAAAACCAGTCTCAGATATCATACTGATTTTTAAGTCAACACTATATATTTTATGATCTTTTCAGTGTGATG GCAAGGTGCTTGTTATGTCTAGAAAGTAAGAAAACAATATGAGGAGACATTCTGTCTTTCAAAAGGTAATGGTAC TAGAATAATTTTATTAGAATAATTTCATACTGAAATTATTTTCTCCCAAATAAAAATTAGATGGAAAAATGTGAAA AATCTATGATGAAGGTAATTAAAGTAGATTATGGCCTTTTTAAGTATTGCAGTCTAAAACTTCAAAAACTAAAAT CATTGTCAAAATTAATATGATTATTAATCAGAATATCAGATATGATTCACTATTTAAACTATGATAAATTATGAT AATATATGAGGAGGCCTCGCTATAGCAAAAATAGTTAAAAATGCTGACATAACACCAAACTTCATTTTTTAAAAAA TCTGTTGTTCCAAATGTGTATAATTTTAAAGTAATTTCTAAAGCAGTTTATTATAATGGTTTGCCTGCTTAAAAG GTATAATTAAACTTCTTTTCTCTTCTACATTGACACACAGAAATGTGTCAATGTAAAGCCAAAACCATCTTCTGT GTTTATGGCCAATCTATTCTCAAAGTTAAAAGTAAAATTGTTTCAGAGTCACAGTTCCCTTTATTTCACATAAGC ATTAAATTGTGTACACTTTCATTACTTTTAAAAATGTAGAAATTCTTCATGAACATAACTCTGCTGAATGTAAAA GAAAATTTTTTTCAAAAATGCTGTTAATGTATACTACTGGTGGTTGATTGGTTTTATTTTATGTAGCTTGACAA TTCAGTGACTTAATATCTATTCCATTTGTATTGTACATAAAATTTTCTAGAAATACACTTTTTTCCAAAGTGTAA TGTAGTTGTCACACTAAAATTTAATTGGATATTGATGAAATCATTGGCCTGGCAAAATAAAACATGTTGAATTCC CC

895/2825 FIGURE 786

MAFRTICVLVGVFICSICVKGSSQPQARVYLTFDELRETKTSEYFSLSHHPLDYRILLMDEDQDRIYVGSKDHIL SLNINNISQEALSVFWPASTIKVEECKMAGKDPTHGCGNFVRVIQTFNRTHLYVCGSGAFSPVCTYLNRGRRSED QVFMIDSKCESGKGRCSFNPNVNTVSVMINEELFSGMYIDFMGTDAAIFRSLTKRNAVRTDQHNSKWLSEPMFVD AHVIPDGTDPNDAKVYFFFKEKLTDNNRSTKQIHSMIARICPNDTGGLRSLVNKWTTFLKARLVCSVTDEDGPET HFDELEDVFLLETDNPRTTLVYGIFTTSSSVFKGSAVCVYHLSDIQTVFNGPFAHKEGPNHQLISYQGRIPYPRP GTCPGGAFTPNMRTTKEFPDDVVTFIRNHPLMYNSIYPIHKRPLIVRIGTDYKYTKIAVDRVNAADGRYHVLFLG TDRGTVQKVVVLPTNNSVSGELILEELEVFKNHAPITTMKISSKKQQLYVSSNEGVSQVSLHRCHIYGTACADCC LARDPYCAWDGHSCSRFYPTGKRRSRRQDVRHGNPLTQCRGFNLKAYRNAAEIVQYGVKNNTTFLECAPKSPQAS IKWLLQKDKDRRKEVKLNERIIATSQGLLIRSVQGSDQGLYHCIATENSFKQTIAKINFKVLDSEMVAVVTDKWS PWTWASSVRALPFHPKDIMGAFSHSEMQMINQYCKDTRQQHQQGDESQKMRGDYGKLKALINSRKSRNRRNQLPE

896/2825 FIGURE **787**

GACGGCCGCCGGAGCCGAGGCCAACTGTCGCCTGGTTGGGCCCGGAAATGGGACGTCGCGCTTTCTCAGGGAG CAGCAAGCCCGGCCGCAAGACCAACCAGCTGCAGTACATGCAGAATGTGGTGGTGAAGACGCTCTGGAAACACCA GTTCGCCTGGCCCTTCTACCAGCCCGTGGACGCAATCAAATTGAACCTGCCGGATTATCATAAAATAATTAAAAA CCCAATGGATATGGGGACTATTAAGAAGAGACTAGAAAATAATTATTATTGGAGTGCAAGCGAATGTATGCAGGA CTTCAACACCATGTTTACAAATTGTTACATTTATAACAAGCCCACAGATGACATAGTGCTAATGGCCCAAGCTTT AGAGAAAATTTTTCTACAAAAAGTGGCCCAGATGCCCCAAGAGGAAGTTGAATTATTACCCCCTGCTCCAAAGGG CACTGCAAACGTCACGTCCCAGTCCCCCCAGCTGCCGCCCCACCTCCTCCTGCCACACCCATCGTCCCCGT GGTCCCTCCTACGCCGCCTGTCGTCAAGAAAAAGGGCGTGAAGCGGAAAGCAGACCACTCCCACGACGTC GGCCATCACTGCCAGCCGGAGTGAGTCGCCCCCGCCGTTGTCAGACCCCAAGCAGGCCAAAGTGGTGGCCCGGCG GGAGAGTGGTGGCCCCCATCAAGCCTCCCAAGAAGGACCTGGAGGACGGCGAGGTGCCCCAGCACGCAGGCAA GAAGGGCAAGCTGTCGGAGCACCTGCGCTACTGCGACAGCATCCTCAGGGAGATGCTATCCAAGAAGCACGCGGC CTACGCCTGGCCCTTCTACAAGCCAGTGGATGCCGAGGCCCTGGAGCTGCACGACTACCACGACATCATCAAGCA CCCGATGGACCTCAGCACCGTGAAAAGGAAGATGGATGGCCGAGAGTACCCAGACGCACAGGGCTTTGCTGCTGA TGTCCGGCTGATGTTCTCGAATTGCTACAAATACAATCCCCCAGACCACGAGGTTGTGGCCATGGCCCGGAAGCT GGCCCCCATGGTGAGCAAGGGCGCTGAGAGCAGCCGTAGCAGTGAGGAGAGCTCTTCGGACTCAGGCAGCTCGGA CTCGGAGGAGCGGCCCACCAGGCTGCCGGAGCTGCAGGAGCAGCTGAAGGCCGTGCACGAGCAGCTGGCCGC GGAGAAGGAGAAGGACACAAAGTGAAGGCCGAGGAAGAAGAAGGCCCAAGGTGGCTCCGCCTGCCAAGCA CAAGCAGGCATCTGCCTCCTACGACTCAGAGGAAGAGGAGGGGCCTGCCCATGAGCTACGATGAAAAGCGCCA GCTTAGCCTGGACATCAACCGGCTGCCCGGGGAGAAGCTGGGCCGGGTAGTGCACATCATCCAATCTCGGGAGCC CTCGCTCAGGGACTCCAACCCCGACGAGATAGAAATTGACTTTGAGACTCTGAAACCCACCACTTTGCGGGAACT GGAGAGATATGTCAAGTCTTGTTTACAGAAAAAGCAAAGGAAACCGTTCTCAGCAAGCGGGAAGAAACAGGCAGC CAAGTCGAAAGAGGAGCTAGCTCAGGAAAAGAAGAAGAAGCGTCTGCAGGATGTCAGCGGCCAGCT GAGCAGCAGCAAGAAGCCCGCCCGGAAAGAAGCCCGGCTCAGCACCCTCAGGGGGCCCGTCCAGGCTCAGCAG CAGCAGCTCCTCCGAGTCTGGGAGCAGCAGCTCCAGCGGGTCCAGCTCTGACAGCAGTGACTCAGAA**TGA**ACTGG CTTCGGACAGAACAGGACAGATGTCGCACACGCCGAGACTCTGCCGTACCCCTCTGTGGTTCATATTACTA CTTCTGTTCCATGGTGTGCAGGTCTGCTTCTTAATTCAGTGTTATGATATCTTCCAGTTTTTTGCTTTCATAGGTC AGAGATCTATCTTGTGTGTGCGTTAGACTTGATGAGAAGGTGTGAACTCTGCAGAAAGTCTCTTCTTCATCACTG AATTCAGTCACTTGGAGATGACAACTTCAAATGCTAACCCGATGACCCCAGAAAACCGTGTGAGATTCGTACCGA CTCGGATATGGGTACTGTATTTGAAAGCTGTTAGCTTTGTCAACACGCATTGTCCTTGTCATTTGGGCCCCGAGC GCTGGGCCTAAGCGAAGGGTCATCGCAGCCCAGCCCGGAGCGTGGAGCCCTTGGGGGGTGGTCGGGTGGGATGT GCGTTCTCCGCTCGTGGTGATGTCAGGAGCTCCTCGGAGGGGAACAGAGCGGCTGTGTATGCAGCCTGCAGGTTTC **GGCATCTATG**

897/2825 FIGURE **788**

MSTATTVAPAGIPATPGPVNPPPPEVSNPSKPGRKTNQLQYMQNVVVKTLWKHQFAWPFYQPVDAIKLNLPDYHK IIKNPMDMGTIKKRLENNYYWSASECMQDFNTMFTNCYIYNKPTDDIVLMAQALEKIFLQKVAQMPQEEVELLPP APKGKGRKPAAGAQSAGTQQVAAVSSVSPATPFQSVPPTVSQTPVIAATPVPTITANVTSVPVPPAAAPPPPATP IVPVVPPTPPVVKKKGVKRKADTTTPTTSAITASRSESPPPLSDPKQAKVVARRESGGRPIKPPKKDLEDGEVPQ HAGKKGKLSEHLRYCDSILREMLSKKHAAYAWPFYKPVDAEALELHDYHDIIKHPMDLSTVKRKMDGREYPDAQG FAADVRLMFSNCYKYNPPDHEVVAMARKLQDVFEMRFAKMPDEPVEAPALPAPAAPMVSKGAESSRSSEESSSDS GSSDSEEERATRLAELQEQLKAVHEQLAALSQAPVNKPKKKKEKKEKKKKKDKEKEKEKHKVKAEEEKKAKVAP PAKQAQQKKAPAKKANSTTTAGRQLKKGGKQASASYDSEEEEEGLPMSYDEKRQLSLDINRLPGEKLGRVVHIIQ SREPSLRDSNPDEIEIDFETLKPTTLRELERYVKSCLQKKQRKPFSASGKKQAAKSKEELAQEKKKELEKRLQDV SGQLSSSKKPARKEKPGSAPSGGPSRLSSSSSSSSSSSSSSSSSSS

898/2825 FIGURE 789A

GAATTCCGGGAGCCCGGAGCTGCTGCAGCAGCCCGAACCGGCGTCCGAGTCCGGGGAAGGCCCCCGCGAGCG GAGGGCCCCTGGGGGTGCAAGAGGCGAATTGGCTGCCCGGGGTGCAGGAGGGACAGGTGGTGCGGGCTGTCAGG GAGGAGGAGGTGGTGACTGGCAGGCCCCGGGCCCCACGGTACCTCCGGGGCTGAAGGGGACGCAGGATGTAGGGGC ATGGGGAGTCGGGCGCAGAAGAGTGCGGGGAACGCTGAGCTCTGGGAGCCACTGCCTGAGGGCAGGCCGCGGCCG GCGGGAACCTCTTCTGCCGTCTCAGCCTGGGCGTCGCTGAAGCTGTGTCTGCGGGGAGGCAGCGGAAGGCGGCAG AGGCTGGGAGGCGGCAGAATCCCAGAGGAGGGGCACAGGCTGCTGCTGGAGCTGCTGCTCCGAGGTGCAGCTG CAACAGTACTTCCTGCGGCTCCGAGATGACCTCAACGTCACCCGCCTGTCCCACTTTGAGTACGTCAAGAATGAG GACCTGGAGAAGATCGGCATGGGTCGGCCTGGCCAGCGGCGGCTGTGGGGAGGCTGTGAAGAGGAGGAGGAGGCCTTG TGCAAACGCAAGTCGTGGATGAGTAAGGTGTTCAGTGGAAAGCGACTGGAGGCTGAGTTCCCACCTCATCACTCT CAGAGCACCTTCCGGAAGACCTCGCCCGCCCCTGGGGGCCCAGCAGGGGGAGGGCCCCTGCAGAGCCTCACCTGC CTCATTGGGGAGAAGGACCTGCGCCTCCTGGAGAAGCTGGGTGATGGTTCCTTTGGCGTGGTGCGCAGGGGCGAG GTGGTGCTCACGCCCCCATGAAGATGGTGACAGAGCTGGCACCTCTGGGATCGTTGTTGGACCGGCTACGTAAG CACCAGGGCCACTTCCTCCTGGGGACTCTGAGCCGCTACGCTGTGCAGGTGGCTGAGGGCATGGGCTACCTGGAG TCCAAGCGCTTTATTCACCGTGACCTGGCTGCCCGCAATCTGCTGTTGGCTACCCGCGACCTGGTCAAGATCGGG GACTTTGGGCTGATGCGAGCACTACCTCAGAATGACGACCATTACGTCATGCAGGAACATCGCAAGGTGCCCTTC GCCTGGTGTGCCCCCGAGAGCCTGAAGACACGCACCTTCTCCCATGCCAGCGACACCTGGATGTTCGGGGTGACA CTGTGGGAAATGTTCACCTACGGCCAGGAGCCCTGGATCGGCCTCAACGGCAGTCAGATCCTGCATAAGATCGAC AAGGAGGGGGAGCCCCCGGCCCGAGGACTGTCCCCAGGACATCTACAACGTCATGGTCCAGTGCTGGGCT CACAAGCCAGAGGACAGACCCACGTTTGTGGCCCTGCGGGACTTCCTGCTGGAGGCCCAGCCCACAGACATGCGG TCCGTGGCCGGCCTGTCGGCCCAGGACATCAGCCAGCCCCTGCAGAACAGCTTCATCCACACAGGGCATGGCGAC AGTGACCCCCGCCACTGCTGGGGCTTCCCGGACAGGATTGACGAACTGTATCTGGGAAACCCCATGGACCCCCC CCTGTGAGCGAGGACCAAGACCCCTTGTCCAGCGACTTCAAGAGGCTGGGCCTGCGGAAGCCAGGCCTGCCCCGA GGGCTGTGGCTGGCGAAGCCCTCGGCGCGGGTGCCGGGCACCAAGGCCAGCCGAGGCAGCGGGGCTGAGGTCACG GACGCCTGCTCCCTGCTGGACGAGACCCCGCCTCAGAGCCCCACGCGGCACTGCCCCGGCCCCTGCACCCCACG TACGCCTTTGTGCCTGAGCAGGCGCCGCCCCCCCCCTGGAGGACAACCTGTTCCTCCCGCCCCAGGGTGGG GGCAAGCCGCCCAGCTCCGCACAGACCGCAGAGATCTTCCAGGCGCTACAGCAGGAGTGCATGAGGCAACTGCAG GCTCCGGGCTCCCCGGCCCCTCTCCCAGCCCGGGGGGTGACGACAAGCCCCAGGTGCCTCCTCGGGTACCCATC GGACCTGCTTCCCCTCCCGGGGTGCCTCCGCGGGAGCCCCTGTCCCCTCAAGGCTCGAGGACACCCAGCCCCTG GTACCACCTGGCAGCTCCCCGCTGCCACCCCGGCTCTCAAGCTCACCTGGGAAGACCATGCCCACCACCACAGAGC TTTGCCTCAGACCCCAAGTACGCCACCCCCAGGTGATCCAGGCCCCTGGCGCGGGTGGTCCCTGCATCCTGCCC ATCGTCCGGGATGGCAAGAAGGTCAGCAGCACCCACTATTACTTGCTGCCCGAGCGACCATCCTACCTGGAGCGC CCCAGCACCCCAGCCCCCGCCGCCCCCCCACGGCCACCGTGCGGCCGATGCCCCAGGCTGCCTTGGACCCCAAGGCC AACTTCTCCACCAACAACAGCAACCCAGGGGCCCGGCCACCCCCGAGGGCCACTGCTCGGCTGCCACAGAGG GGCTGCCCTGGCGATGGGCCAGAGGCGGGCCGGCCAGCAGACAAGATCCAGATGGCCATGGTGCATGGGGTGACC ACAGAGGAGTGCCAGGCCCTGCAGTGCCACGGCTGGAGCGTGCAGAGGGCTGCCCAGTATCTGAAGGTGGAG

899/2825 FIGURE 789B

900/2825 FIGURE **790**

MQPEEGTGWLLELLSEVQLQQYFLRLRDDLNVTRLSHFEYVKNEDLEKIGMGRPGQRRLWEAVKRRKALCKRKSW
MSKVFSGKRLEAEFPPHHSQSTFRKTSPAPGGPAGEGPLQSLTCLIGEKDLRLLEKLGDGSFGVVRRGEWDAPSG
KTVSVAVKCLKPDVLSQPEAMDDFIREVNAMHSLDHRNLIRLYGVVLTPPMKMVTELAPLGSLLDRLRKHQGHFL
LGTLSRYAVQVAEGMGYLESKRFIHRDLAARNLLLATRDLVKIGDFGLMRALPQNDDHYVMQEHRKVPFAWCAPE
SLKTRTFSHASDTWMFGVTLWEMFTYGQEPWIGLNGSQILHKIDKEGERLPRPEDCPQDIYNVMVQCWAHKPEDR
PTFVALRDFLLEAQPTDMRALQDFEEPDKLHIQMNDVITVIEGRAENYWWRGQNTRTLCVGPFPRNVVTSVAGLS
AQDISQPLQNSFIHTGHGDSDPRHCWGFPDRIDELYLGNPMDPPDLLSVELSTSRPPQHLGGVKKPTYDPVSEDQ
DPLSSDFKRLGLRKPGLPRGLWLAKPSARVPGTKASRGSGAEVTLIDFGEEPVVPALRPCPPSLAQLAMDACSLL
DETPPQSPTRALPRPLHPTPVVDWDARPLPPPPAYDDVAQDEDDFEICSINSTLVGAGVPAGPSQGQTNYAFVPE
QARPPPPLEDNLFLPPQGGGKPPSSAQTAEIFQALQQECMRQLQAPGSPAPSPSPGGDDKPQVPPRVPIPPRPTR
PHVQLSPAPPGEEETSQWPGPASPPRVPPREPLSPQGSRTPSPLVPPGSSPLPPRLSSSPGKTMPTTQSFASDPK
YATPQVIQAPGAGGPCILPIVRDGKKVSSTHYYLLPERPSYLERYQRFLREAQSPEEPTPLPVPLLLPPPSTPAP
AAPTATVRPMPQAALDPKANFSTNNSNPGARPPPPRATARLPQRGCPGDGPEAGRPADKIQMAMVHGVTTEECQA
ALQCHGWSVQRAAQYLKVEQLFGLGLRPRGECHKVLEMFDWNLEQAGCHLLGSWGPAHHKR

901/2825 FIGURE **791A**

CAGTCGCTCCGAGCGGCCGCGAGCAGAGCCGCCCAGCCCTGTCAGCTGCGCCGGGACGATAAGGAGTCAGGCCAG CTATTCCAGAGAGGACAACTGTCATCTGGGAAGTAACAACGCAGG**ATG**CCCCCTGGGGTGGACTGCCCCATGGAA TTCTGGACCAAGGAGGAGAATCAGAGCGTTGTGGTTGACTTCCTGCTGCCCACAGGGGTCTACCTGAACTTCCCT GTGTCCCGCAATGCCAACCTCAGCACCATCAAGCAGCTGCTGTGGCACCGCGCCCAGTATGAGCCGCTCTTCCAC ATGCTCAGTGGCCCCGAGGCCTATGTGTTCACCTGCATCAACCAGACAGCGGAGCAGCAAGAGCTGGAGGACGAG CAACGCCTCTGTGTGACGTGCACCCTTCCTGCCCCTCCTGCGCCTGGTGGCCCGTGAGGGCGACCGCGTGAAG AAGCTCATCAACTCACAGATCAGCCTCCTCATCGGCAAAGGCCTCCACGAGTTTGACTCCTTGTGCGACCCAGAA GCCTGGCTGCAGTACAGTTTCCCCCTGCAGCTGGAGCCCTCGGCTCAAACCTGGGGGCCTGGTACCCTGCGGCTC $\tt CCGAACCGGGCCCTTCTGGTCAACGTTAAGTTTGAGGGCAGCGAGGAGAGCTTCACCTTCCAGGTGTCCACCAAG$ GACGTGCCGCTGGCGCTGATGCCCTGCGGAAGAAGGCCACAGTGTTCCGGCAGCCGCTGGTGGAGCAG CCGGAAGACTACACGCTGCAGGTGAACGCCAGGCATGAGTACCTGTATGGCAACTACCCGCTCTGCCAGTTCCAG TACATCTGCAGCTGCCTGCACAGTGGGTTGACCCCTCACCTGACCATGGTCCATTCCTCCTCCATCCTCGCCATG CGGGATGAGCAGAGCAACCCTGCCCCCAGGTCCAGAAACCGCGTGCCAAACCACCTCCCATTCCTGCGAAGAAG CCTTCCTCTGTGTCCCTGTGGTCCCTGGAGCAGCCGTTCCGCATCGAGCTCATCCAGGGCAGCAAAGTGAACGCC GACGAGCGGATGAAGCTGGTGCTGCCAGGCCGGCTTTTCCACGGCAACGAGATGCTGTGCAAGACGGTGTCCAGC TCGGAGGTGAGCGTGTGCTCGGAGCCCGTGTGGAAGCAGCGGCTGGAGTTCGACATCAACATCTGCGACCTGCCC TCCAAGAAGGCGGACTGCCCCATTGCCTGGGCCAACCTCATGCTGTTTGACTACAAGGACCAGCTTAAGACCGGG GAACGCTGCCTCTACATGTGGCCCTCCGTCCCAGATGAGAAGGGCGAGCTGCTGAACCCCACGGGCACTGTGCGC CCCGCCTGGAGAAGATCTTGGAGCTGGGGCGACACAGCGAGTGTGTCACCGAGGAGGAGGAGCAGCTGCAG CTGCGGGAAATCCTGGAGCGGCGGGGTCTGGGGAGCTGTATGAGCACGAGAAGGACCTGGTGTGGAAGCTGCGG CATGAAGTCCAGGAGCACTTCCCGGAGGCGCTAGCCCGGCTGCTGCTGGTCACCAAGTGGAACAAGCATGAGGAT GTGGCCCAGATGCTCTACCTGCTGTGCTCCTGGCCGGAGCTGCCCGTCCTGAGCGCCCTGGAGCTGCTAGACTTC TACCTGCTGCAGCTGGTGCAGGTGCTCAAGTACGAGTCCTACCTGGACTGCGAGCTGACCAAATTCCTGCTGGAC CGGGCCCTGGCCAACCGCAAGATCGGCCACTTCCTTTTCTGGCACCTCCGCTCCGAGATGCACGTGCCGTCGGTG GCCCTGCGCTTCGGCCTCATCCTGGAGGCCTACTGCAGGGGCAGCACCCACACATGAAGGTGCTGATGAAGCAG GGGGAAGCACTGAGCAAACTGAAGGCCCTGAATGACTTCGTCAAGCTGAGCTCTCAGAAGACCCCCAAGCCCCAG GACCCCAGCACCCTGCTGGCTGAAGTCTGCGTGGAGCAGTGCACCTTCATGGACTCCAAGATGAAGCCCCTGTGG ATCATGTACAGCAACGAGGAGGCAGCCGGCGGCAGCGTGGGCATCATCTTTAAGAACGGGGATGACCTCCGG CAGGACATGCTGACCCTGCAGATGATCCAGCTCATGGACGTCCTGTGGAAGCAGGAGGGGCTGGACCTGAGGATG ACCCCTATGGCTGCCTCCCCACCGGGGACCGCACAGGCCTCATTGAGGTGGTACTCCGTTCAGACACCATCGCC AACATCCAACTCAACAAGAGCAACATGGCAGCCACAGCCGCCTTCAACAAGGATGCCCTGCTCAACTGGCTGAAG ${\tt TCCAAGAACCCGGGGGAGGCCCTGGATCGAGCCATTGAGGAGTTCACCCTCTCCTGTGCTGGCTATTGTGTGGCC}$ ACATATGTGCTGGGCATTGGCGATCGGCACAGCGACACATCATGATCCGAGAGAGTGGGCAGCTGTTCCACATT GATTTTGGCCACTTTCTGGGGAATTTCAAGACCAAGTTTGGAATCAACCGCGAGCGTGTCCCATTCATCCTCACC TATGACTTTGTCCATGTGATTCAGCAGGGGAAGACTAATAATAGTGAGAAATTTGAACGGTTCCGGGGCTACTGT CTGCCTGAGCTCAGCTGCTCCAAAGACATCCAGTATCTCAAGGACTCCCTGGCACTGGGGAAAACAGAGGAGGAG CACAACGTGTCCAAAGACAACAGGCAGTAGTGGCTCCTCCCAGCCCTGGGCCCAAGAGGGGGGGCTGCGGGTCG TGGGGACCAAGCACATTGGTCCTAAAGGGGCTGAAGAGCCTGAACTGCACCTAACGGGAAAGAACCGACATGGCT GCCTTTTGTTTACACTGGTTATTTATTATGACTTGAAATAGTTTAAGGAGCTAAACAGCCATAAACGGAAACGC CTCCTTCATTCAGCGGCGGTGCTGGGCCCCCGAGGCTGCACCTGGCTCTCGGCTGAGGATTGTCACCCCAAGTC

902/2825 FIGURE **791B**

TTCCAGCTGGTGGATCTGGGCCCAGCAAAGACTGTTCTCCTCCCGAGGGAACCTTCTTCCCAGGCCTCCCGCCAG ${\tt ACTGCCTGGGTCCTGGCGGTCACCTGGTGCCTACTGTCCGACAGGATGCCTCGATCCTCGTGCGACCCC}$ ACCCTGTGTATCCTCCCTAGACTGAGTTCTGGCAGCTCCCCGAGGCAGCCGGGGTACCCTCTAGATTCAGGGATG CTTGCTCTCCACTTTTCAAGTGGGTCTTGGGTACGAGAATTCCCTCATCTTTCTCTACTGTAAAGTGATTTTGTT TGCAGGTAAGAAAATAATAGATGACTCACCACACCTCTACGGCTGGGGAGATCAGGCCCAGCCCCATAAAGGAGA ATCTACGCTGGTCCTCAGGACGTGTTAAAGAGATCTGGGCCTCATGTAGCTCACCCCGGTCACGCATGAAGGCAA $\tt CTGTTGCCCAGGCTGGAGTGCAGTGGTGCAATCTTGGCTCACTGTAACCTCCGCCTCCCAGGTTCAAGTGATTCT$ TCTTGCCTCAGCCTCCTGAGTAGCTGGGATTACAGGTGTGCACCACCCGTACCCAGCTAATTTTTGTATTTTAGT AGAGACGGGGGTTTCACCATGTTGGCTGGGCTGGTCTCGAACTCCTGACCTCAGGTGATCCACCCGCCTGAGCCT CCCAAAGTGCTGGGATTACAGGCATGAGCCACCACGCCCGGCCCACTCTGCCATTGTCTAAGCCACCTCTGAAAG CAGGTTTTAACAAAAGGATGAGGCCAGAACTCTTCCAGAACCATCACCTTTGGGAACCTGCTGTGAGAGTGCTGA GGTACCAGAAGTGTGAGAACGAGGGGGCGTGCTGGGATCTTTCTCTCTGACTATACTTAGTTTGAAATGGTGCAG GCTTAGTCTTAAGCCTCCAAAGGCCTGGATTTGAGCAGCTTTAGAAATGCAGGTTCTAGGGCCTTCTCCCAGCCTT CAGAAGCCAACTAACTCTGCAGATGGGGCTAGGACTGTGGGCTTTTAGCAGCCCACAGGTGATCCTAACATATCA GGCCATGGACTCAGGACCTGCCCGGTGATGCTGTTGATTTCTCAAAGGTCTTCCAAAACTCAACAGAGCCAGAAG TAGCCGCCCGCTCAGCGCTCAGGTGCCAGCTCTGTTCTGATTCACCAGGGGTCCGTCAGTAGTCATTGCCACCC GCGGGGCACCTCCCTGGCCACACGCCTGTTCCCAGCAAGTGCTGAAACTCACTAGACCGTCTGCCTGTTTCGAAA TGGGGAAAGCCGTGCGCGTTATTTATTTAAGTGCGCCTGTGTGCGCGGGTGTGGGAGCACACTTTGCAAAG $\verb|CCACAGCGTTTCTGGTTTTGGGTGTACAGTCTTGTGTGCCTGGCGAGAAGAATATTTTCTATTTTTTAAGTCAT|\\$ TTCATGTTTCTGTCTGGGGAAGGCAAGTTAGTTAAGTATCACTGATGTGGGTTGAGACCAGCACTCTGTGAAACC

903/2825 FIGURE **792**

MPPGVDCPMEFWTKEENQSVVVDFLLPTGVYLNFPVSRNANLSTIKQLLWHRAQYEPLFHMLSGPEAYVFTCINQ
TAEQQELEDEQRRLCDVQPFLPVLRLVAREGDRVKKLINSQISLLIGKGLHEFDSLCDPEVNDFRAKMCQFCEEA
AARRQQLGWEAWLQYSFPLQLEPSAQTWGPGTLRLPNRALLVNVKFEGSEESFTFQVSTKDVPLALMACALRKKA
TVFRQPLVEQPEDYTLQVNGRHEYLYGNYPLCQFQYICSCLHSGLTPHLTMVHSSSILAMRDEQSNPAPQVQKPR
AKPPPIPAKKPSSVSLWSLEQPFRIELIQGSKVNADERMKLVVQAGLFHGNEMLCKTVSSSEVSVCSEPVWKQRL
EFDINICDLPRMARLCFALYAVIEKAKKARSTKKKSKKADCPIAWANLMLFDYKDQLKTGERCLYMWPSVPDEKG
ELLNPTGTVRSNPNTDSAAALLICLPEVAPHPVYYPALEKILELGRHSECVHVTEEEQLQLREILERRGSGELYE
HEKDLVWKLRHEVQEHFPEALARLLLVTKWNKHEDVAQMLYLLCSWPELPVLSALELLDFSFPDCHVGSFAIKSL
RKLTDDELFQYLLQLVQVLKYESYLDCELTKFLLDRALANRKIGHFLFWHLRSEMHVPSVALRFGLILEAYCRGS
THHMKVLMKQGEALSKLKALNDFVKLSSQKTPKPQTKELMHLCMRQEAYLEALSHLQSPLDPSTLLAEVCVEQCT
FMDSKMKPLWIMYSNEEAGSGGSVGIIFKNGDDLRQDMLTLQMIQLMDVLWKQEGLDLRMTPYGCLPTGDRTGLI
EVVLRSDTIANIQLNKSNMAATAAFNKDALLNWLKSKNPGEALDRAIEEFTLSCAGYCVATYVLGIGDRHSDNIM
IRESGQLFHIDFGHFLGNFKTKFGINRERVPFILTYDFVHVIQQGKTNNSEKFERFRGYCERAYTILRRHGLLFL
HLFALMRAAGLPELSCSKDIQYLKDSLALGKTEEEALKHFRVKFNEALRESWKTKVNWLAHNVSKDNRQ

904/2825 FIGURE **793**

CGCTCCTCGGGCTGCCCTCGGTTGACAATCGTCTCCAGGATGGTCTCTACCATGCTATCTGGCCTACTGTTTTG GCTGGCATCTGGATGGACTCCAGCATTTGCTTACAGCCCCCGGACCCCTGACCGGGTCTCAGAAGCAGATATCCA GAGGCTGCTTCATGGTGTTATGGAGCAATTGGGCATTGCCAGGCCCCGAGTGGAATATCCAGCTCACCAGGCCAT GAATCTTGTGGGCCCCCAGAGCATTGAAGGTGGAGCTCATGAAGGACTTCAGCATTTGGGTCCTTTTGGCAACAT $\tt CCCCAACATCGTGGCAGAGTTGACTGGAGACAACATTCCTAAGGACTTTAGTGAGGATCAGGGGTACCCAGACCC$ TCCAAATCCCTGTCCTGTTGGAAAAACAGATGATGGATGTCTAGAAAACACCCCTGACACTGCAGAGTTCAGTCG AGAGTTCCAGTTGCACCAGCATCTCTTTGATCCGGAACATGACTATCCAGGCTTGGGCAAGTGGAACAAGAAACT CCTTTACGAGAAGATGAAGGGAGAGAGAGACGAAAGCGGAGGAGTGTCAATCCATATCTACAAGGACAGAGACT GGATAATGTTGTTGCAAAGAAGTCTGTCCCCCATTTTTCAGATGAGGATAAGGATCCAGAG**TAA**AGAAAGATGC TAGACGAAAACCCACATTACCTGTTAGGCCTCAGCATGGCTTATGTGCACGTGTAAATGGAGTCCCTGTGAATGA ATTCTGCTTTTTGCTAAATTAGAATAAGAGCTTTTTTGTTTCTTGGGTTTTTAAAATGTGAATCTGCAATGATCA TAAAAATTAAAATGTGAATGTCAACAATAAAAAGCAAGACTATGAAAGGCTCAGATTTCTTGCAGTTTAAAATGG TGTCTGAGGTTGTACTATTTTGGCCAAGTCTGTAGAAAGCTGTCATTTGATTTTGATTATGTAGTTCATCCAGCC CTTGGGCATTGTTATACACCAGTAAAGAAGGCTGTACTCAAGAGGAGGAGCTGACACATTTCACTTGGCTGCGTC TTAATAAACATGAATGCAAGCATTGGC

905/2825 FIGURE **794**

MVSRMVSTMLSGLLFWLASGWTPAFAYSPRTPDRVSEADIQRLLHGVMEQLGIARPRVEYPAHQAMNLVGPQSIE GGAHEGLQHLGPFGNIPNIVAELTGDNIPKDFSEDQGYPDPPNPCPVGKTDDGCLENTPDTAEFSREFQLHQHLF DPEHDYPGLGKWNKKLLYEKMKGGERRKRRSVNPYLQGQRLDNVVAKKSVPHFSDEDKDPE

906/2825 FIGURE **795**

907/2825 FIGURE **796**

 ${\tt MLLPNILLTGTPGVGKTTLGKELASKSGLKYINVGDLAREEQLYDGYDEEYDCPILDEDRVVDELDNQMREGGVI} \\ {\tt VDYHGCDFFPERWFHIVFVLRTDTNVLYERLETRGYNEKKLTDNIQCEIFQVLYEEATASYKEEIVHQLPSNKPE} \\ {\tt ELENNVDQILKWIEQWIKDHNS} \\$

908/2825 FIGURE **797**

CAGTGCCTTGGTAATGACCAGGGCTCCAGAAAGAGAGTGTCCTTGTGGCTGGGGGGCCCCTGTGCCTGACATTCCTCCTGACTCTGCGGTGGAGCTGTGGAAGCCAGGCGCACAGGATGCAAGCAGCCAGGCCCAGGGAGGCAGCAGCTGCA TCCTCAGAGAGGAAGCCAGGATGCCCCACTCTGCTGGGGGTACTGCAGGGGTGGGGGCTGGAGGCTGCAGAGCCCA CAGCCCTGCTCACCAGGGCAGAGCCCCCTTCAGAACCCACAGAGATCCGTCCACAAAAGCGGAAAAAGGGGCCAG CCCCCAAAATGCTGGGGAACGAGCTATGCAGCGTGTGTGGGGACAAGGCCTCGGGCTTCCACTACAATGTTCTGA GCTGCGAGGGCTGCAAGGGATTCTTCCGCCGCAGCGTCATCAAGGGAGCGCACTACATCTGCCACAGTGGCGGCC ACTGCCCCATGGACACCTACATGCGTCGCAAGTGCCAGGATGTCGGCTTCGCAAATGCCGTCAGGCTGGCATGC GGGAGGAGTGTCCTGTCAGAAGAACAGATCCGCCTGAAGAAACTGAAGCGGCAAGAGGAGGAACAGGCTCATG CCACATCCTTGCCCCCAGGCGTTCCTCACCCCCCAAATCCTGCCCCAGCTCAGCCCGGAACAACTGGGCATGA TCGAGAAGCTCGTCGCCCAGCAACAGTGTAACCGGCGCTCCTTTTCTGACCGGCTTCGAGTCACGCCTTGGC CCATGGCACCAGATCCCCATAGCCGGGAGGCCCGTCAGCAGCGCTTTGCCCACTTCACTGAGCTGGCCATCGTCT CTGTGCAGGAGATAGTTGACTTTGCTAAACAGCTACCCGGCTTCCTGCAGCTCAGCCGGGAGGACCAGATTGCCC TGCTGAAGACCTCTGCGATCGAGGTGATGCTTCTGGAGACATCTCGGAGGTACAACCCTGGGAGTGAGAGTATCA CCTTCCTCAAGGATTTCAGTTATAACCGGGAAGACTTTGCCAAAGCAGGGCTGCAAGTGGAATTCATCAACCCCA TCTTCGAGTTCTCCAGGGCCATGAATGAGCTGCAACTCAATGATGCCGAGTTTGCCTTGCTCATTGCTATCAGCA TCTTCTCTGCAGACCGGCCCAACGTGCAGGACCAGCTCCAGGTGGAGAGGCTGCAGCACACATATGTGGAAGCCC TGCATGCCTACGTCTCCATCCACCATGACCGACTGATGTTCCCACGGATGCTAATGAAACTGGTGAGCC TCCGGACCCTGAGCAGCGTCCACTCAGAGCAAGTGTTTGCACTGCGTCTGCAGGACAAAAAGCTCCCACCGCTGC CCTGGTGGCTGCCTCCTAGAAGTGGAACAGACTGAGAAGGGCAAACATTCCTGGGAGCTGGGCAAGGAGATCCTC CCGTGGCATTAAAAGAGAGTCAAAGGGT

909/2825 FIGURE 798

MSLWLGAPVPDIPPDSAVELWKPGAQDASSQAQGGSSCILREEARMPHSAGGTAGVGLEAAEPTALLTRAEPPSE PTEIRPQKRKKGPAPKMLGNELCSVCGDKASGFHYNVLSCEGCKGFFRRSVIKGAHYICHSGGHCPMDTYMRRKC QECRLRKCRQAGMREECVLSEEQIRLKKLKRQEEEQAHATSLPPRRSSPPQILPQLSPEQLGMIEKLVAAQQQCN RRSFSDRLRVTPWPMAPDPHSREARQQRFAHFTELAIVSVQEIVDFAKQLPGFLQLSREDQIALLKTSAIEVMLL ETSRRYNPGSESITFLKDFSYNREDFAKAGLQVEFINPIFEFSRAMNELQLNDAEFALLIAISIFSADRPNVQDQ LQVERLQHTYVEALHAYVSIHHPHDRLMFPRMLMKLVSLRTLSSVHSEQVFALRLQDKKLPPLLSEIWDVHE

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910/2825 FIGURE **799**

CACCCTGCACCTGCAGACGGGGAACCTGCTGAACTGGGGCCGCCTGCGGAAGAAGTGCCCGTCCACGCACAGCGA GGAGCTTCATGATTGTATCCAAAAAACCTTGAATGAATGGAGTTCCCAAATCAACCCAGATTTGGTCAGGGAGTT TCCAGATGTCTTGGAATGCACTGTATCTCATGCAGTAGAAAAGATAAATCCTGATGAAAGAGAAAATGAAAGT TTCTGCAAAACTGTTCATTGTAGAATCAAACTCTTCATCATCAACTAGAAGTGCAGTTGACATGGCCTGTTCAGT CCTTGGAGTTGCACAGCTGGATTCTGTGATCATTGCTTCACCTCCTATTGAAGATGGAGTTAATCTTTCCTTGGA GCATTTACAGCCTTACTGGGAGGAATTAGAAAACTTAGTTCAGAGCAAAAAGATTGTTGCCATAGGTACCTCTGA ${\tt TCTAGACAAAACAGTTGGAACAGCTGTATCAGTGGGCACAGGTAAAACCAAATAGTAACCAAGTTAATCTTGC}$ CTCCTGCTGTGTGATGCCACCAGATTTGACTGCATTTGCTAAACAATTTGACATACAGCTGTTGACTCACAATGA TCCAAAAGAACTGCTTTCTGAAGCAAGTTTCCAAGAAGCTCTTCAGGAAAGCATTCCTGACATTCAAGCGCACGA $\tt GTGGGTGCCGCTGTGGCTACTGCGGTATTCGGTCATTGTGAAAAGTAGAGGAATTATCAAATCAAAAGGCTACAT$ AAATTGAGATGTGTAAAATCTAGTTACTGCCTGTAAATGGTGTCATTGAGGCCAGATATTCTTTCGTCATATTTGA CAGTATGTTGTCTGTCAAGTTTTAAATACTTATCTTGCCTCCATATCAATCCATTCTCATGAACCTCTGTATTGC TTTCCTTAAACTATTGTTTTCTAATTGAAATTGTCTATAAAGAAAATACTTGCAATATATTTTTCCTTTATTTTT ATGACTAATATAAATCAAGAAAATTTGTTGTTAGATATATTTTGGCCTAGGTATCAGGGTAATGTATATACATAT TTTTTATTCCAAAAAAATTCATTAATTGCTTCTTAACTCTTATTATAACCAAGCAATTTAATTACAATTGTTA AAACTGAAATACTGGAAGAAGATATTTTTCCTGTCATTGATGAGATATATCAGAGTAACTGGAGTAGCTGGGATT TACTAGTAGTGTAAATAAAATTCACTCTTCAATAC

911/2825 FIGURE 800

MGTDSRAAKALLARARTLHLQTGNLLNWGRLRKKCPSTHSEELHDCIQKTLNEWSSQINPDLVREFPDVLECTVS HAVEKINPDEREEMKVSAKLFIVESNSSSSTRSAVDMACSVLGVAQLDSVIIASPPIEDGVNLSLEHLQPYWEEL ENLVQSKKIVAIGTSDLDKTQLEQLYQWAQVKPNSNQVNLASCCVMPPDLTAFAKQFDIQLLTHNDPKELLSEAS FQEALQESIPDIQAHEWVPLWLLRYSVIVKSRGIIKSKGYILQAKRRGS

912/2825 FIGURE 801

 $\texttt{AGACACCTCTGCCCTCACC} \underline{\textbf{ATG}} \\ \texttt{AGCCTCTGGCAGCCCCTGGTCCTGGTGCTCCTGGTGCTGCTTTGC} \\$ ${\tt TGCCCCCAGACAGCGCCAGTCCACCCTTGTGCTCTTCCCTGGAGACCTGAGAACCAATCTCACCGACAGGCAGCT}$ GGCAGAGGAATACCTGTACCGCTATGGTTACACTCGGGTGGCAGAGATGCGTGGAGAGTCGAAATCTCTGGGGCC TGCGCTGCTTCTCCAGAAGCAACTGTCCCTGCCCGAGACCGGTGAGCTGGATAGCGCCACGCTGAAGGCCAT GCGAACCCCACGGTGCGGGGTCCCAGACCTGGGCAGATTCCAAACCTTTGAGGGCGACCTCAAGTGGCACCACCA $\verb|CTTCGCACTGTGGAGCGCGGTGACGCCGCTCACCTTCACTCGCGTGTACAGCCGGGACGCAGACATCGTCATCCA| \\$ GTTTGGTGTCGCGGAGCCGGGGTATCCCTTCGACGGGAAGGACGGCCTCCTGGCACACGCCTTTCCTCC TGGCCCCGGCATTCAGGAGACGCCCATTTCGACGATGACGAGTTGTGGTCCCTGGGCAAGGGCGTCGTGGTTCC AACTCGGTTTGGAAACGCAGATGGCGGGCCTGCCACTTCCCCTTCATCTTCGAGGGCCGCTCCTACTCTGCCTG CACCACCGACGGTCGCTCCGACGGCTTGCCCTGGTGCAGTACCACGGCCAACTACGACACCGACGACCGGTTTGG CTTCTGCCCCAGCGAGAGACTCTACACCCGGGACGGCAATGCTGATGGGAAACCCTGCCAGTTTCCATTCATCTT CCAAGGCCAATCCTACTCCGCCTGCACCACGGACGGTCGCTCCGACGGCTACCGCTGGTGCGCCACCACCGCCAA CTACGACCGGGACAAGCTCTTCGGCTTCTGCCCGACCCGAGCTGACTCGACGGTGATGGGGGGCAACTCGGCGGG GGAGCTGTGCGTCTTCCCCTTCACTTTCCTGGGTAAGGAGTACTCGACCTGTACCAGCGAGGGCCGCGGAGATGG GCGCCTCTGGTGCGCTACCACCTCGAACTTTGACAGCGACAAGAAGTGGGGCTTCTGCCCGGACCAAGGATACAG TTTGTTCCTCGTGGCGGCGCATGAGTTCGGCCACGCGCTTAGATCATTCCTCAGTGCCGGAGGCGCTCAT GTACCCTATGTACCGCTTCACTGAGGGGCCCCCCTTGCATAAGGACGACGTGAATGGCATCCGGCACCTCTATGG TCCTCGCCCTGAACCTGAGCCACGGCCTCCAACCACCACCACCACGCCACGGCTCCCCCGACGGTCTGCCC CACCGGACCCCCACTGTCCACCCCTCAGAGCGCCCCACAGCTGGCCCCACAGGTCCCCCCTCAGCTGGCCCCAC AGGTCCCCCCACTGCTGGCCCTTCTACGGCCACTACTGTGCCTTTGAGTCCGGTGGACGATGCCTGCAACGTGAA CATCTTCGACGCCATCGCGGAGATTGGGAACCAGCTGTATTTGTTCAAGGATGGGAAGTACTGGCGATTCTCTGA GGGCAGGGGGAGCCGCAGGGCCCCTTCCTTATCGCCGACAAGTGGCCCGCGCTGCCCCGCAAGCTGGACTC GGTCTTTGAGGAGCCGCTCTCCAAGAAGCTTTTCTTCTTCTCTGGGCGCCAGGTGTGGGTGTACACAGGCGCGTC GGTGCTGGGCCCGAGGCGTCTGGACAAGCTGGGCCTGGGAGCCGACGTGGCCCAGGTGACCGGGGCCCTCCGGAG TGGCAGGGGAAGATGCTGCTGTTCAGCGGGCGCCCCTCTGGAGGTTCGACGTGAAGGCGCAGATGGTGGATCC GGGCTACGTGACCTATGACATCCTGCAGTGCCCTGAGGACTACGGCTCCCGTCCTGCTTTGCAGTGCCATGTAAA TCCCCACTGGGACCAACCCTGGGGAAGGAGCCAGTTTGCCGGATACAAACTGGTATTCTGTTCTGGAGGAAAGGG CTAACCTTT

913/2825 FIGURE 802

MSLWQPLVLVLLVLGCCFAAPRQRQSTLVLFPGDLRTNLTDRQLAEEYLYRYGYTRVAEMRGESKSLGPALLLLQ KQLSLPETGELDSATLKAMRTPRCGVPDLGRFQTFEGDLKWHHHNITYWIQNYSEDLPRAVIDDAFARAFALWSA VTPLTFTRVYSRDADIVIQFGVAEHGDGYPFDGKDGLLAHAFPPGPGIQGDAHFDDDELWSLGKGVVVPTRFGNA DGAACHFPFIFEGRSYSACTTDGRSDGLPWCSTTANYDTDDRFGFCPSERLYTRDGNADGKPCQFPFIFQGQSYS ACTTDGRSDGYRWCATTANYDRDKLFGFCPTRADSTVMGGNSAGELCVFPFTFLGKEYSTCTSEGRGDGRLWCAT TSNFDSDKKWGFCPDQGYSLFLVAAHEFGHALGLDHSSVPEALMYPMYRFTEGPPLHKDDVNGIRHLYGPRPEPE PRPPTTTTPQPTAPPTVCPTGPPTVHPSERPTAGPTGPPSAGPTGPPTAGPSTATTVPLSPVDDACNVNIFDAIA EIGNQLYLFKDGKYWRFSEGRGSRPQGPFLIADKWPALPRKLDSVFEEPLSKKLFFFSGRQVWVYTGASVLGPRR LDKLGLGADVAQVTGALRSGRGKMLLFSGRRLWRFDVKAQMVDPRSASEVDRMFPGVPLDTHDVFQYREKAYFCQ DRFYWRVSSRSELNQVDQVGYVTYDILQCPED

914/2825 FIGURE 803

GCGGTGGTGGACACGTCGAGCCGGGTAGAAGTGGAGGGGCCGTTCGAAGAGTCGTGAGGGGGTGACGGGTTAAGA TTCGGAGAGAGGTGCTAGTGGCTGGACTTGACCTGGAAAGAATCTTCTGCTGACTCTCAACTTTTCCTGGAAA AAATGGATCATTCCCACCATATGGGGATGAGCTATATGGACTCCAACAGTACCATGCAACCTTCTCACCATCACC CAACCACTTCAGCCTCACACTCCCATGGTGGAGGAGACAGCAGCATGATGATGATGCCTATGACCTTCTACTTTG GCTTTAAGAATGTGGAACTACTGTTTTCCGGTTTTGGTGATCAATACAGCTGGAGAAATGGCTGGAGCTTTTGTGG GCATTCGCTACAATTCCATGCCTGTCCCAGGACCAAATGGAACCATCCTTATGGAGACACACAAAACTGTTGGGC AACAGATGCTGAGCTTTCCTCACCTCCTGCAAACAGTGCTGCACATCATCCAGGTGGTCATAAGCTACTTCCTCA TGCTCATCTTCATGACCTACAACGGGTACCTCTGCATTGCAGTAGCAGCAGGGGCCGGTACAGGATACTTCCTCT TCAGCTGGAAGAAGGCAGTGGTAGTGGATATCACAGAGCATTGCCAT**TGA**CATCAAACTCTATGGCGTGGCCTTA TCGATTGCAGTGGGAAGTTGTTGAAGACTTGAAGACGTGATTCCTGCTCCAATCATCCCTTCTTGCTCCTCTTTG TGCACGTACACACACACACACACACACACACACACACCCCTGCTCAACAGAGGTTTAGTTTACAGTCTC TGAACTAAAGTAGTAACCTCCCAAATTGTTTTTTCTAATAAGCTGAGATTCCCATTTCTCTTAAGGAGAAGCCAC $\verb|CCATGAGATGTCTTTTCCTTCTCCATCATCTTAGAGCCAAGTTATATGTTCTTGTCTAATCCATGTAGCTTTTTG|\\$ GGTCAGTGATGGAAAGGGGTTAACTTCAGCCAGGATTGATGGCAGCTGAGGGAAATTCTTGCCCAACTAAACCCA GAACTCAAACTTAACATTAGAAAATAAGGTCCAGGGCCGGACACAGTGGCCCATGCCTGTAATCCCAGCACTTTG GGGGGCCAAGGCAGGCTGGATCACCTGAGGACAGGAGTTCGAGACCAGTCTGGCCAACATGGGGAAACCCCGTCT CTACTAAAAATACAAAAATTAGCCGGGCATGGTGGTGGGCGCCTGTAATCCCAGCTACTCAGAAGGCTGAGGCAG GAGAATCACTTGAACCTAGGAGGCGGAGGTTGCAGTGAGCCAAGATGGCGCCATTGCACTCCAGCCTGGGTGACA AGAGTGAAACTCCATCTCAAAAAAAAAAAGAAAGGTCCAGCTTTTGGATTCAATGAGTGGGAAATACATTGTG CCTTTCTCTAGATGTGATACGTTATACCAAAATCTTTGTAGTGTGCAGAGCGGTGGTTTGAGACTAAATACAGGC $\tt TTAGAACTTGCAGAGTGTATTCTTGGATGGCTGATGCATCGACTTGCATTCCCACTTAACACTTTGATTAGCC$ GTTT

915/2825 FIGURE 804

 ${\tt MDHSHHMGMSYMDSNSTMQPSHHHPTTSASHSHGGGDSSMMMMPMTFYFGFKNVELLFSGLVINTAGEMAGAFVA} \\ {\tt VFLLAMFYEGLKIARESLLRKSQVSIRYNSMPVPGPNGTILMETHKTVGQQMLSFPHLLQTVLHIIQVVISYFLMLFMTYNGYLCIAVAAGAGTGYFLFSWKKAVVVDITEHCH} \\ {\tt LIFMTYNGYLCIAVAAGAGTGYFLFSWKKAVVVDITEHCH} \\ {\tt MISSING MARKET MARKE$

916/2825 FIGURE 805

GCGCGCCCCTGCCCTGCGGAGCCTGCGCCCTTCTACGAACCGGGCCGGGCGGCAAGCCGGGCCGCGGGC CGCCTACATCGACTCCATGGCCGCCGTGCCCACCCTGGAGCTGTGCCACGACGAGCTCTTCGCCGACCTCTTCAA CCCTGCCGCTCCCGCCTGCTCAAGCGCGAGCCCGACTGGGGCGACGCGACGCGCCCGGCTCGCTGTTGCCCGC GGACCGCGGCAGCCCCGAGTACCGGCAGCGCGCGCGAGACAACATCGCCGTGCGCAAGAGCCGCGACAAGGC CAAGCGGCGCAACCAGGAGATGCAGCAGAAGTTGGTGGAGCTGTCGGCTGAGAACGAGAAGCTGCACCAGCGCGT CGCCGGGACAGCAGACTGCCGG<u>TAA</u>CGCGCGGCCGGGCGGGAGAGACTCAGCAACGACCCATACCTCAGACCCG ACGGCCCGGAGCGGAGCGCCCTGCCCTGGCGCAGCCAGAGCCGCCGGGTGCCCGCTGCAGTTTCTTGGGACAT AGGAGCGCAAAGAAGCTACAGCCTGGACTTACCACCACTAAACTGCGAGAAAGCTAAACGTGTTTATTTTCCCT TATTAATTTTTACAGTATTTGTAAGAATAAAGCAGCATTTGAAATCGC

917/2825 FIGURE 806

 ${\tt MSAALFSLDGPARGAPWPAEPAPFYEPGRAGKPGRGAEPGALGEPGAAAPAMYDDESAIDFSAYIDSMAAVPTLE} \\ {\tt LCHDELFADLFNSNHKAGGAGPLELLPGGPARPLGPGPAAPRLLKREPDWGDGDAPGSLLPAQVGPCAQTVVSLA} \\ {\tt AAGQPTPPTSPEPPRSSPRQTPAPGPAREKSAGKRGPDRGSPEYRQRRERNNIAVRKSRDKAKRRNQEMQQKLVE} \\ {\tt LSAENEKLHQRVEQLTRDLAGLRQFFKQLPSPPFLPAAGTADCR} \\$

918/2825 FIGURE 807

AGGGGTCTAGCTGGCCTTTGCTCGGCCCTCCCCAGCGCCCGGCTTTGAACCCGCCCTGCACTGCTGTCTGGGCCG CTGAGGTGGGCGTGCGAGGGCCGGCCGTGGCCTCTGCCCCCACGGGGCCAGAGCCAAGGCCGATCCCTGCC GCCCTCCCCTCGGACAAGGCCACCGGAGCTCCCGGAGCCGGGCCTGGTGTCCGGCGGCGGCAACGGAGCTTAGAG GCCAGTGCCCCGCTCTTGGAGCAAGTGATGCGGCAAGAGGGAAAGTACCCAGTACGGAACGACATGGAGCTATGG AAGGAGCACCGGGACCAGCACCTGACCTATGGGCCGTTCACCACGGAAGGACACCACTGGTACCAGCTGCGC GACTTTATGACTCGACTGGACCAGCTGCGGGCAGAGAGTGCTTCGGGGAACCAGGTGTCGGACATGGCTCAACTC ATGGCTGGAGTGGACACGACATCCAACACGCTGACATGGGCCCTGTACCACCTCTCAAAGGACCCTGAGATCCAG GAGGCCTTGCACGAGGAAGTGGTGGGTGGTGCCAGCCGGGCAAGTGCCCCAGCACAAGGACTTTGCCCACATG CCGTTGCTCAAAGCTGTGCTTAAGGAGACTCTGCGTCTCTACCCTGTGGTCCCCACAAACTCCCGGATCATAGAA AAGGAAATTGAAGTTGATGGCTTCCTCTTCCCCAAGAACACCCAGTTTGTGTTCTGCCACTATGTGGTGTCCCGG CTGGAGATGCAGCTACTCCTCGCAAGGCTGATCCAGAAGTACAAGGTGGTCCTGGCCCCGGAGACGGGGGAGTTG AAGAGTGTGGCCCGCATTGTCCTGGTTCCCAATAAGAAAGTGGGCCTGCAGTTCCTGCAGAGACAGTGC**TGA**GCT GAGTCTCCGCCTTGCTGGGGCTTGTCCTAGAGGCTCCAGCTCTGGCACAGTGGTTCCTGGCTGCCATGTCTC AGATGAGGAGGAGAGAGGAGGCCGCCAGACTCGAGAGGTGGGAGGAACTCCTTGCACACACCCTGAGCTTTTG AATATAAAATAAAGGGACTTTTATTTCTTATTGG

919/2825 FIGURE 808

MAALGCARLRWALRGAGRGLCPHGARAKAAIPAALPSDKATGAPGAGPGVRRRQRSLEEIPRLGQLRFFFQLFVQGYALQLHQLQVLYKAKYGPMWMSYLGPQMHVNLASAPLLEQVMRQEGKYPVRNDMELWKEHRDQHDLTYGPFTTEGHWYQLRQALNQRLLKPAEAALYTDAFNEVIDDFMTRLDQLRAESASGNQVSDMAQLFYYFALEAICYILFEKRIGCLQRSIPEDTVTFVRSIGLMFQNSLYATFLPKWTRPVLPFWKRYLDGWNAIFSFGKKLIDEKLEDMEAQLQAAGPDGIQVSGYLHFLLASGQLSPREAMGSLPELLMAGVDTTSNTLTWALYHLSKDPEIQEALHEEVVGVVPAGQVPQHKDFAHMPLLKAVLKETLRLYPVVPTNSRIIEKEIEVDGFLFPKNTQFVFCHYVVSRDPTAFSEPESFQPHRWLRNSQPATPRIQHPFGSVPFGYGVRACLGRRIAELEMQLLLARLIQKYKVVLAPETGELKSVARIVLVPNKKVGLQFLQRQC

920/2825 FIGURE 809

921/2825 FIGURE 810

 ${\tt MCDAFVGTWKLVSSENFDDYMKEVGVGFATRKVAGMAKPNMIISVNGDVITIKSESTFKNTEISFILGQEFDEVTADDRKVKSTITLDGGVLVHVQKWDGKSTTIKRKREDDKLVVECVMKGVTSTRVYERA}$

922/2825 FIGURE 811

GTAAGGCCGTTTTCTTTTCCCATTCGCTCATCTGCCAGGAAAAGGGACTTGCCGTTGGCGCTTCGGCCTCTTGTT CATTGAGAAAAAGAGGAAATACTCCGCGTGCGCTTGTAGAAGGGGAGTCGTCTCCAGCTCCGAACCCCGGAGTG TTCATCAGCGGGGAATCTGGCTCCGAATTCTCTTTTTTTCTCCCGCCGATTGCTCGGAAGTTGGTCTAAAGCAGA GAGAGGGGTTGGTGCAAAGCCGCGATCACGGAGTTCAGATGTGTTCTAAGCCTGCTGGAGTGACCACACTTCCAA GACCTGATGGAGGCCAGAGCTCAGAGTGGCAACGGGTCGCAGCCCTTGCTGCAGACGCCCCGTGACGGTGGCAGA CAGCGTGGGGAGCCCGACCCCAGAGACGCCCTCACCCAGCAGGTACATGTCTTGTCTCTGGATCAGATCAGAGCC ATCCGAAACACCAATGAGTACACAGAGGGGCCTACTGTCGTCCCAAGACCTGGGCTCAAGCCTGCTCCTCGCCCC TCCACTCAGCACAAACACGAGAGACTCCACGGTCTGCCTGAGCACCGCCAGCCTCCTAGGCTCCAGCACTCGCAG GTCCATTCTTCTGCACGAGCCCCTCTGTCCAGATCCATAAGCACGGTCAGCTCAGGGTCGCGGAGCAGTACGAGG ACAAGTACCAGCAGCAGCTCCTCTGAACAGAGACTGCTAGGATCATCCTTCTCCTCCGGGCCTGTTGCTGATGGC ATAATCCGGGTGCAACCCAAATCTGAGCTCAAGCCAGGTGAGCTTAAGCCACTGAGCAAGGAAGATTTGGGCCTG CACGCCTACAGGTGTGAGGACTGTGGCAAGTGCAAATGTAAGGAGTGCACCTACCCAAGGCCTCTGCCATCAGAC TGGATCTGCGACAAGCAGTGCCTTTGCTCGGCCCAGAACGTGATTGACTATGGGACTTGTGTATGCTGTGTAAA GGTCTCTTCTATCACTGTTCTAATGATGATGAGGACAACTGTGCTGACAACCCATGTTCTTGCAGCCAGTCTCAC TGTTGTACACGATGGTCAGCCATGGGTGTCATGTCCCTCTTTTTTGCCTTGTTTATGGTGTTACCTTCCAGCCAAG GGTTGCCTTAAATTGTGCCAGGGGTGTTATGACCGGGTTAACAGGCCTGGTTGCCGCTGTAAAAAACTCAAACACA GTTTGCTGCAAAGTTCCCACTGTCCCCCCTAGGAACTTTGAAAAAACCAACA**TAG**CATCATTAATCAGGAATATTA TAATAGAAAGGTGGGATAGTCTTTGCTGTTTGCGGTGAAATGCTTTTTGTCCATGTGCCGTTTTAACTGATATGC TTGTTAGAACTCAGCTAATGGAGCTCAAAGTATGAGATACAGAACTTGGTGACCCATGTATTGCATAAGCTAAAG CAACACAGACACTCCTAGGCAAAGTTTTTGTTTGTGAATAGTACTTGCAAAACTTGTAAATTAGCAGATGACTTT TTTCCATTGTTTTCTCCAGAGAGAATGTGCTATATTTTTGTATATACAATAATATTTTGCAACTGTGAAAAACAAG AGTATGTTTTTAGATTGTATTTTGCCTTACAGAAAGCCTTTATTGTAAGACTCTGATTTCCCTTTGGACTTCATG TATATTGTACAGTTACAGTAAAATTCAACCTTTATTTTCTAATTTTTTCAACATATTGTTTAGTGTAAAGAATAT GAGGATGTGATAGTTGCTGCAAATGAGGGGTTACAGATGCATATGTCCAATATAAAATAGAAAATATATTAACGT TTGAAATTAAAAAAAAAAAAAAAAAA

923/2825 FIGURE 812

$$\label{thm:condition} \begin{align} MEARAQSGNGSQPLLQTPRDGGRQRGEPDPRDALTQQVHVLSLDQIRAIRNTNEYTEGPTVVPRPGLKPAPRPST\\ QHKHERLHGLPEHRQPPRLQHSQVHSSARAPLSRSISTVSSGSRSSTRTSTSSSSSEQRLLGSSFSSGPVADGII\\ RVQPKSELKPGELKPLSKEDLGLHAYRCEDCGKCKCKECTYPRPLPSDWICDKQCLCSAQNVIDYGTCVCCVKGL\\ FYHCSNDDEDNCADNPCSCSQSHCCTRWSAMGVMSLFLPCLWCYLPAKGCLKLCQGCYDRVNRPGCRCKNSNTVC\\ CKVPTVPPRNFEKPT\\ \end{align*}$$

924/2825 FIGURE 813

GGAGGTCCAGAGATCCTGAGAGCCCCGAGCGCCCCAGCGCCCCGGGAGAGTTGAGAATT**ATG**GCTGCAGGAAAA TTTGCAAGCCTTCCCAGAAACATGCCGGTGAATCACCAGTTCCCCCTGGCCTCATCCATGGACCTTCTGAGCAGC AGGTCCCCTCTCGCTGAGCATCGCCCAGATGCCTATCAAGATGTCTATACATGGCACCCTTCCACGGAAGAAA AAAGGTCCTCCCCATAAGGTCCTGTGATGACTTCAGTCACATGGGCACCCTCCCCACTCCAAATCCCCACGG CAGAACTCGCCTGTGACCCAGGATGGCATCCAGGAGAGCCCATGGCAGGACCGGCACGGCGAAACCTTCACCTTC AGGGATCCACATCTTCTGGACCCAACTGTGGAATATGTGAAGTTCTCCAAGGAGAGGCACATCATGGACAGGACC CCCGAGAAACTGAAGAAGGAGCTGGAGGAGGAGCTGCTCCTGAGCAGCGAGGACCTGCGCAGCCATGCCTGGTAC CACGGCCGCATCCCCGACAGGTGTCTGAAAACCTTGTGCAGCGAGATGGTGACTTCCTAGTTCGTGACTCTCTG TCCAGCCCTGGGAACTTTGTCCTGACCTGTCAGTGGAAGAACCTCGCTCAGCACTTCAAAATCAACCGGACAGTT CTGCGACTCAGCGAGGCCTACAGCCGCGTGCAGTACCAGTTCGAGATGGAGAGCTTCGACTCCATCCCCGGCCTG GTGCGCTGCTACGTGGGCAACCGCCGGCCCATCTCCCAGCAGAGTGGCGCCATCATCTTCCAGCCCATCAACAGG ACGGTGCCTCTGCGGTGCCTGGAGGAGCATTATGGCACCTCCCCAGGCCAGGCCCGGGAGGGCAGCCTCACCAAG GGAAGGCCGGATGTGGCCAAGAGGCTGAGCCTCACCATGGGTGGCGTCCAGGCCCGAGAGCAGAATTTGCCCAGG TTGTCCCTCAAAGCCCACCAGTCAGAGAGCTACCTGCCGATTGGCTGCAAGCTGCCACCTCAGTCCTCGGGTGTG GACACAAGCCCCTGCCCAAACTCACCTGTGTTCAGGACGGGAAGCGAGCCTGCCCTGAGCCCAGCAGTGGTTCGG AGGGTCTCCTCAGACGCCAGGGCTGGGGAGGCGCTGAGGGGATCAGACAGTCAACTGTGCCCTAAGCCCCCGCCT AAGCCCTGCAAGGTGCCGTTCCTCAAGGTTCCCTCGTCTCCCTCTGCCTGGCTCAACTCAGAGGCCAACTACTGT GAACTGAACCCAGCGTTTGCCACAGGCTGCGGCAGGGGAGCCAAAGCTACCCTCATGTGCCCCAGGGAAGCCACACA GAACTGCTCACAGCCAAGCAGAATGAGGCGCCAGGTCCCCGGAACTCTGGCGTCAACTACTTGATCCTTGATGAT GATGACAGGGAAAGACCTTGGGAACCTGCGGCAGCTCAGATGGAGAAGGGGCAGTGGGACAAGGGCGAGTTTGTG CCCCTGGAAACAGCAATGTTGAAACGTGCAAAAGAACTGTTCACCAACGACCCCCAAGGTCATCGCCCAGCAC GTACTGAGCATGGACTGCAGGGTTGCTAGGATACTTGGAGTCTCTGAAGAGATGAGGAGGAACATGGGGGTGAGC TCAGGCCTGGAACTCATTACCTTGCCTCACGGACACCAGCTGCGCCTGGACATAATTGAAAGACACAACAATG GCCATCGGCATTGCAGTGGACATTCTGGGATGCACGGGCACTTTGGAGGACCGAGCGGCCACTCTGAGTAAGATC ATCCAGGTGGCGGTGGAACTGAAGGATTCCATGGGGGACCTCTATTCCTTCTCAGCTCTCATGAAAGCCCTGGAA ATGCCACAGATCACAAGGTTAGAAAAGACGTGGACTGCTCTGCGGCACCAGTACACCCAAACTGCCATTCTCTAT GAGAAACAGCTGAAGCCCTTCAGCAAACTCCTGCATGAAGGCAGAGAGTCCACATGTGTTCCCCCAAACAATGTA TCAGTCCCACTGCTGATGCCGCTTGTGACGTTAATGGAGCGCCAGGCTGTGACTTTTGAAGGAACCGACATGTGG GAAAAAAACGACCAGAGCTGTGAAATCATGCTGAACCATTTGGCAACAGCGCGATTCATGGCCGAGGCTGCAGAC GAACCTTTAGAATATCTTTTCAAGTTTCCCCAGCTTCATCTTTTGGGAAAGCTTACTGTTTTTGATAAAGTAATAA TGTGCAAATCTGACAATATACAAGCTTTTAGTATCCACAGGATATTAAACGTGTAAATTGCACAGAGCACACTTA TTTATGAATTGTCTAAAGTTACTACTGATTTTAAAATGAATAATTTATTATTAAAGGTAACTACTGCTAATGTTGA TCAGCAAATTTAAGAGAAGACCTAGCTATGTTGGCTGGTTGCTTTCTATTATCATGGTATTTGACCATTTTAGTT TTAATTCCATGTCAGATAAGTGTAAATAGAAGAGTTTAAAAGCATGAAACATTTCAGAAGGTATCAGTTATATGA AAAA

925/2825 FIGURE 814

MAAGKFASLPRNMPVNHQFPLASSMDLLSSRSPLAEHRPDAYQDVSIHGTLPRKKKGPPPIRSCDDFSHMGTLPH
SKSPRQNSPVTQDGIQESPWQDRHGETFTFRDPHLLDPTVEYVKFSKERHIMDRTPEKLKKELEEELLLSSEDLR
SHAWYHGRIPRQVSENLVQRDGDFLVRDSLSSPGNFVLTCQWKNLAQHFKINRTVLRLSEAYSRVQYQFEMESFD
SIPGLVRCYVGNRRPISQQSGAIIFQPINRTVPLRCLEEHYGTSPGQAREGSLTKGRPDVAKRLSLTMGGVQARE
QNLPRGNLLRNKEKSGSQPACLDHMQDRRALSLKAHQSESYLPIGCKLPPQSSGVDTSPCPNSPVFRTGSEPALS
PAVVRRVSSDARAGEALRGSDSQLCPKPPPKPCKVPFLKVPSSPSAWLNSEANYCELNPAFATGCGRGAKLPSCA
QGSHTELLTAKQNEAPGPRNSGVNYLILDDDDRERPWEPAAAQMEKGQWDKGEFVTPLLETVSSFRPNEFESKFL
PPENKPLETAMLKRAKELFTNNDPKVIAQHVLSMDCRVARILGVSEEMRRNMGVSSGLELITLPHGHQLRLDIIE
RHNTMAIGIAVDILGCTGTLEDRAATLSKIIQVAVELKDSMGDLYSFSALMKALEMPQITRLEKTWTALRHQYTQ
TAILYEKQLKPFSKLLHEGRESTCVPPNNVSVPLLMPLVTLMERQAVTFEGTDMWEKNDQSCEIMLNHLATARFM
AEAADSYRMNAERILAGFQPDEEMNEICKTEFQMRLLWGSKGAQVNQTERYEKFNQILTALSRKLEPPPVKQAEL

926/2825 FIGURE 815

CCTCCCCTCGCCCGGCGCGCTCCGCCTCTCGCTCGCCTCCCCCTCCGCTCTCCGAGGCGCCCGG AGGATACGCGCTCGGCGCTGGGACGCGACTGCGCTCAGTTCTCTCCTCTCGGAAGCTGCAGCCATGATGGAAGTT GCCCTCTCAGCGCCTGTGAGCAGCCGCGGGGGCAGCCCCTCGGGGAGCCGGCCTGCGGCGGCGGCAGCG GCGGCGTTTCTCGCCTCCTCTTCGTCTTTTCTAACCGTGCAGCCTCTTCCTCGGCTTCTCCTGAAAGGGAAGGTG TCCAGGGACCCGGGCCGGTTTTAAACCTCCCGTCCGCCGCCGCACCCCCGTGGCCCGGGCTCCGGAGGCCG $\verb|CCGGCGGAGGCAGCCGTTCGGAGGATTATTCGTCTTCTCCCCATTCCGCTGCCGCCGCTGCCAGGCCTCTGGCTG|\\$ CTGAGGAGAGCAGCCCAGTCGCTGCAACCATCCAGCAGCCGCCGCAGCAGCCATTACCCGGCTGCGGTCCAGA GCCAAGCGGCGCAGAGCGAGGGGCATCAGCTACCGCCAAGTCCAGAGCCATTTCCATCCTGCAGAAGAAGCCCC ACATTATTGCTATGGGATTTCCTGCAGAAAGACTTGAAGGCGTATACAGGAACAATATTGATGATGTAAGGT TTTTGGATTCAAAGCATAAAAACCATTACAAGATATACAATCTTTGTGCTGAAAGACATTATGACACCGCCAAAT TTAATTGCAGAGTTGCACAATATCCTTTTGAAGACCATAACCCACACAGCTAGAACTTATCAAACCCTTTTGTG AAGATCTTGACCAATGGCTAAGTGAAGATGACAATCATGTTGCAGCAATTCACTGTAAAGCTGGAAAGGGACGAA $\verb|CTGGTGTAATGATATGTGCATATTTATTACATCGGGGCAAATTTTTAAAGGCACAAGAGGCCCTAGATTTCTATG|\\$ GGGAAGTAAGGACCAGAGACAAAAAGGGAGTAACTATTCCCAGTCAGAGGCGCTATGTGTATTATTATAGCTACC TGTTAAAGAATCATCTGGATTATAGACCAGTGGCACTGTTGTTTCACAAGATGATGTTTGAAACTATTCCAATGT TCAGTGGCGGAACTTGCAATCCTCAGTTTGTGGTCTGCCAGCTAAAGGTGAAGATATATTCCTCCAATTCAGGAC AGTTCTTCCACAAACAGAACAAGATGCTAAAAAAGGACAAAATGTTTCACTTTTGGGTAAATACATTCTTCATAC CAGGACCAGAGGAAAACCTCAGAAAAAGTAGAAAATGGAAGTCTATGTGATCAAGAAATCGATAGCATTTGCAGTA ACAAAGCCAACCGATACTTTTCTCCAAATTTTAAGGTGAAGCTGTACTTCACAAAAACAGTAGAGGAGCCGTCAA ${\tt ATCCAGAGGCTAGCAGTTCAACTTCTGTAACACCAGATGTTAGTGACAATGAACCTGATCATTATAGATATTCTG}$ ACACCACTGACTCTGATCCAGAGAATGAACCTTTTGATGAAGATCAGCATACAAAATTACAAAAGTC**TGA**TTT GCAATAGGACATTGTGTCAGATTACCAGTTATAGGAACAATTCTCTTTTCCTGACCAATCTTGTTTTACCCTATA CATCCACAGGGTTTTGACACTTGTTGTCCAGTTGAAAAAAAGGTTGTGTAGCTGTGTCATGTATATACCTTTTTGT GTCAAAAGGACATTTAAAATTCAATTAGGATTAATAAAGATGGCACTTTCCCGTTTTATTCCAGTTTTATAAAAA GTGGAGACAGACTGATGTGTATACGTAGGAATTTTTTCCTTTTGTGTTCTGTCACCAACTGAAGTGGCTAAAGAG CTTTGTGATATACTGGTTCACATCCTACCCCTTTGCACTTGTGGCAACAGATAAGTTTGCAGTTGGCTAAGAGAG GTTTCCGAAAGGTTTTGCTACCATTCTAATGCATGTATTCGGGTTAGGGCAATGGAGGGGAATGCTCAGAAAGGA GTGCTAGAAAAGGCAGCTAAAGGAAGTGAATCTGTATTGGGGTACAGGAATGAACCTTCTGCAACATCTTAAGAT CCACAAATGAAGGGATATAAAAATAATGTCATAGGTAAGAAACACAGCAACAATGACTTAACCATATAAATGTGG AGGCTATCAACAAAGAATGGGCTTGAAACATTATAAAAATTGACAATGATTTATTAAATATGTTTTCTCAATTGT AAAAAAAAA

927/2825 FIGURE 816

MTAIIKEIVSRNKRRYQEDGFDLDLTYIYPNIIAMGFPAERLEGVYRNNIDDVVRFLDSKHKNHYKIYNLCAERH YDTAKFNCRVAQYPFEDHNPPQLELIKPFCEDLDQWLSEDDNHVAAIHCKAGKGRTGVMICAYLLHRGKFLKAQE ALDFYGEVRTRDKKGVTIPSQRRYVYYYSYLLKNHLDYRPVALLFHKMMFETIPMFSGGTCNPQFVVCQLKVKIY SSNSGPTRREDKFMYFEFPQPLPVCGDIKVEFFHKQNKMLKKDKMFHFWVNTFFIPGPEETSEKVENGSLCDQEI DSICSIERADNDKEYLVLTLTKNDLDKANKDKANRYFSPNFKVKLYFTKTVEEPSNPEASSSTSVTPDVSDNEPD HYRYSDTTDSDPENEPFDEDQHTQITKV

928/2825 FIGURE 817

AGCGGCCGCCGCCACAGCTGCAGTCAGCACCGTCACCCCAGCAGCATCCGCGCGCCTGCACCGCGCGTGCGG ATCAGCGCGCTACCTGCTGACACGGAACCCTCACCTCAACAAGGACTTGGCCTTTACCCTGGAAGAGACAGC AATTGAACATTCATGGATTGTTGCCACCTTCCTTCAACAGTCAGGAGATCCAGGTTCTTAGAGTAGAAAAATT TCGAGCATCTGAACTCTGACTTTGACAGGTATCTTCTCTTAATGGATCTCCAAGATAGAAAATGAAAAACTCTTTT ATAGAGTGCTGACATCTGACATTGAGAAATTCATGCCTATTGTTTATACTCCCACTGTGGGTCTGGCTTGCCAAC AATATAGTTTGGTGTTTCGGAAACCAAGAGGTCTCTTTATTACTATCCACGATCGAGGGCATATTGCTTCAGTTC TCAATGCATGGCCAGAAGATGTCATCAAGGCCATTGTGGTGACTGATGGAGAGCGTATTCTTGGCTTGGGAGACC TTGGCTGTAATGGAATGGCCATCCCTGTGGGTAAATTGGCTCTATATACAGCTTGCGGAGGGATGAATCCTCAAG AATGTCTGCCTGTCATTCTGGATGTGGGAACCGAAAATGAGGAGTTACTTAAAGATCCACTCTACATTGGACTAC GGCAGAGAAGAGTAAGAGGTTCTGAATATGATGATTTTTTTGGACGAATTCATGGAGGCAGTTTCTTCCAAGTATG GCATGAATTGCCTTATTCAGTTTGAAGATTTTGCCAATGTGAATGCATTTCGTCTCCTGAACAAGTATCGAAACC AGTATTGCACATTCAATGATATTCAAGGAACAGCATCTGTTGCAGGTCTCCCTTGCAGCTCTTCGAA TAACCAAGAACTGTCTGATCAAACAATACTATTCCAAGGAGCTGGGGAGGCTGCCCTAGGGATTGCACACC TGATTGTGATGGCCTTGGAAAAAGAAGGTTTACCAAAAGAGAAAGCCATCAAAAAGATATGGCTGGTTGATTCAA AAGGATTAATAGTTAAGGGACGTGCTTCCTTAACACAAGAGAAAGAGAAGTTTGCCCATGAACATGAAGAAATGA AGAACCTAGAAGCCATTGTTCAAGAAATAAAACCAACTGCCCTCATAGGAGTTGCTGCAATTGGTGGTGCATTCT CAGAACAAATTCTCAAAGATATGGCTGCCTTCAATGAACGGCCTATTATTTTTTGCTTTGAGTAATCCAACTAGCA AAGCAGAATGTTCTGCAGAGCAGTGCTACAAAATAACCAAGGGACGTGCAATTTTTGCCAGTGGCAGTCCTTTTG ATCCAGTCACTCTCCAAATGGACAGACCCTATATCCTGGCCAAGGCAACAATTCCTATGTGTTCCCTGGAGTTG CTCTTGGTGTTGTGGCGTGTGGATTGAGGCAGATCACAGATAATATTTTCCTCACTACTGCTGAGGTTATAGCTC AGCAAGTGTCAGATAAACACTTGGAAGAGGGTCGGCTTTATCCTCCTTTGAATACCATTAGAGATGTTTCTCTGA AAATTGCAGAAAAGATTGTGAAAGATGCATACCAAGAAAAGACAGCCACAGTTTATCCTGAACCGCAAAACAAAG AAGCATTTGTCCGCTCCCAGATGTATAGTACTGATTATGACCAGATTCTACCTGATTGTTATTCTTGGCCTGAAG AGGTGCAGAAAATACAGACCAAAGTTGACCAG**TAG**GATAATAGCAAACATTTCTAACTCTATTAATGAGGTCTTT AAACCTTTCATAATTTTTAAAGGTTGGAATCTTTTATAATGATTCATAAGACACTTAGATTAAGATTTTACTTTA ACAGTCTAAAAATTGATAGAAGAATATCGATATAAATTGGGATAAACATCACATGAGACAATTTTGCTTCACTTT GAGAAACTCATCATTTTTATACAGGACACTAATGGGAAGACCAAAATTACTAATAAATTGACATAACCAACATTA GCGTAGGTGAACTTTTGCTAAGCAGAAATAACACTACTTTGTTGCCTAGAGAAAGATAACTTCTCAAGTATTTTT ATTCCAGTCCTAGATCATATATGTTCTTTTGTGCAACGGAATTCTAACAGTTCTAAGAGAAAGATCACTGCTGTT TACAGCGCCTTGTGCAGCCTTAGATTTTAATATTCTTTTGTCATTGTTACATCTCATAGAGTAAAGCTCTTATTA CCTTGATCCTGAGTCAGAAATCCCACCTGAAATCACCTTTTTTCCCCCTTGATCAAACATCCCATCCTTCAGCTA CCATACTGTTGCTACAGGGATTTTGTGGACTGTGGCCCCTGTCCCGAGGTTGGCACCTTCAGTTCAGCACAGCCT GAGCAGTGAGAAGGTCTGAAAGGAGAGTATATAGTTAAGATCCTTGAGAAAGGGCTGCCTGAGGAACTGACCTCT CTCAGGTAATGAAGAGCTAACTGTGTTATAATCATCTTGCTTTTGCCTGAATTTGGAGAAAGTATTATAATTAAG TTCCCAGTATCAGAAATGTCCTTACATAAGATTAAAATATCTTGATGACTAATACCATTCTATGAGAAAGAGTAG TTATATGCCCAGACTGTATTAATTTACTTTAGAAACTAATGTTTGAAGTAATGGAAAAAATTTTAAATTATAAAG CTAAGGTGCAATAACATTTGCTACTTATTTATAGAATTATTTGAAGAATTTTTGTTTTTTGAAGTAATGCTTTAAGG AGTATAAGATATTCAAGATAAATTATACTATAAAATGATTTTATTGAAAGTTGAAGGTTACACAAATTGTTTTAG GTATGAGCAGAAGAGGTTAAGGTATTTCTAAAGGTAACATATAGTCAAGAGTTTCCTCAAAATAGTTATTTGGAG AAGAATCAGAATGTCTGTGTATTTCTTGTCTGTTTCTATGTTGTCTATAGCTCTGACTAAATGTGTTTACCTAT GCAAAAGATTTATTAAAGCATAGAAAAGGTGAATGAATAACACCCTAAACCCCCGAAGAGAAAAAGAAATCCCG

929/2825 FIGURE 818

RPPPPQLQSAPSPQQHPRACTARAARPGLTPPPNPAPAMEPEAPRRRHTHQRGYLLTRNPHLNKDLAFTLEERQQ LNIHGLLPPSFNSQEIQVLRVVKNFEHLNSDFDRYLLLMDLQDRNEKLFYRVLTSDIEKFMPIVYTPTVGLACQQ YSLVFRKPRGLFITIHDRGHIASVLNAWPEDVIKAIVVTDGERILGLGDLGCNGMGIPVGKLALYTACGGMNPQE CLPVILDVGTENEELLKDPLYIGLRQRRVRGSEYDDFLDEFMEAVSSKYGMNCLIQFEDFANVNAFRLINKYRNQ YCTFNDDIQGTASVAVAGLLAALRITKNKLSDQTILFQGAGEAALGIAHLIVMALEKEGLPKEKAIKKIWLVDSK GLIVKGRASLTQEKEKFAHEHEEMKNLEAIVQEIKPTALIGVAAIGGAFSEQILKDMAAFNERPIIFALSNPTSK AECSAEQCYKITKGRAIFASGSPFDPVTLPNGQTLYPGQGNNSYVFPGVALGVVACGLRQITDNIFLTTAEVIAQ QVSDKHLEEGRLYPPLNTIRDVSLKIAEKIVKDAYQEKTATVYPEPQNKEAFVRSQMYSTDYDQILPDCYSWPEE VQKIQTKVDQ

930/2825 FIGURE 819

GCCTGACCCGCCGCCGAACCCGGCGCCAGCCATGGAGCCCGAAGCCCCCGTCGCCGCCACACCCATCAGCGCG GCTACCTGCTGACACGGAACCCTCACCTCAACAAGGACTTGGCCTTTACCCTGGAAGAGAGACAGCAATTGAACA TTCATGGATTGTTGCCACCTTCCTTCAACAGTCAGGAGATCCAGGTTCTTAGAGTAGAAAAAATTTCGAGCATC TGAACTCTGACTTTGACAGGTATCTTCTCTTAATGGATCTCCAAGATAGAAATGAAAAACTCTTTTATAGAGTGC TGACATCTGACATTGAGAAATTCATGCCTATTGTTTATACTCCCACTGTGGGTCTGGCTTGCCAACAATATAGTT TGGTGTTTCGGAAGCCAAGAGGTCTCTTTATTACTATCCACGATCGAGGGCATATTGCTTCAGTTCTCAATGCAT GGCCAGAAGATGTCATCAAGGCCATTGTGGTGACTGATGGAGAGCGTATTCTTGGCTTGGGAGACCTTGGCTGTA ATGGAATGGGCATCCCTGTGGGTAAATTGGCTCTATATACAGCTTGCGGAGGGATGAATCCTCAAGAATGTCTGC CTGTCATTCTGGATGTGGGAACCGAAATGAGGAGTTACTTAAAGATCCACTCTACATTGGACTACGGCAGAGAA GAGTAAGAGGTTCTGAATATGATGATTTTTTTGGACGAATTCATGGAGGCAGTTTCTTCCAAGTATGGCATGAATT GCCTTATTCAGTTTGAAGATTTTGCCAATGTGAATGCATTTCGTCTCCTGAACAAGTATCGAAACCAGTATTGCA CATTCAATGATGATATTCAAGGAACAGCATCTGTTGCAGTTGCAGGTCTCCTTGCAGCTCTTCGAATAACCAAGA ACAAACTGTCTGATCAAACAATACTATTCCAAGGAGCTGGAGAGGCTGCCCTAGGGATTGCACACCTGATTGTGA TGGCCTTGGAAAAAGAGTTTACCAAAAGAGAAAGCCATCAAAAAGATATGGCTGGTTGATTCAAAAGGATTAA TAGTTAAGGGACGTGCTTCCTTAACACAAGAGAAAGAGAAGTTTGCCCATGAACATGAAGAAATGAAGAACCTAG AAGCCATTGTTCAAGAAATAAAACCAACTGCCCTCATAGGAGTTGCTGCAATTGGTGGTGCATTCTCAGAACAAA TTCTCAAAGATATGGCTGCCTTCAATGAACGGCCTATTATTTTTGCTTTGAGTAATCCAACTAGCAAAGCAGAAT GTTCTGCAGAGCAGTGCTACAAAATAACCAAGGGACGTGCAATTTTTGCCAGTGGCAGTCCTTTTGATCCAGTCA CTCTTCCAAATGGACAGACCCTATATCCTGGCCAAGGCAACAATTCCTACGTGTTCCCTGGAGTTGCTCTTGGTG TTGTGGCGTGTGGATTGAGGCAGATCACAGATAATATTTTCCTCACTACTGCTGAGGTTATAGCTCAGCAAGTGT CAGATAAACACTTGGAAGAGGGTCGGCTTTATCCTCCTTTGAATACCATTAGAGATGTTTCTCTGAAAATTGCAG AAAAGATTGTGAAAGATGCATACCAAGAAAAGACAGCCACAGTTTATCCTGAACCGCAAAACAAAGAAGCATTTG TCCGCTCCCAGATGTATAGTACTGATTATGACCAGATTCTACCTGATTGTTATTCTTGGCCTGAAGAGGTGCAGA AAATACAGACCAAAGTTGACCAG**TAG**GATAATAGCAAACATTTCTAACTCTATTAATGAGGTCTTTAAACCTTTC ATAATTTTTAAAGGTTGGAATCTTTTATAATGATTCATAAGACACTTAGATTAAGATTTTACTTTAACAGTCTAA AAATTGATAGAAGAATATCGATATAAATTGGGATAAACATCACATGAGACAATTTTGCTTCACTTTGCCTTCTGG ATCATTTTTATACAGGACACTAATGGGAAGACCAAAATTACTAATAAATTGAAATAACCAACATTAAAACTCATA ATTATTTTGTTGACCATTTTGTTAAAATCTACTTTTC

931/2825 FIGURE 820

MEPEAPRRHTHQRGYLLTRNPHLNKDLAFTLEERQQLNIHGLLPPSFNSQEIQVLRVVKNFEHLNSDFDRYLLL MDLQDRNEKLFYRVLTSDIEKFMPIVYTPTVGLACQQYSLVFRKPRGLFITIHDRGHIASVLNAWPEDVIKAIVV TDGERILGLGDLGCNGMGIPVGKLALYTACGGMNPQECLPVILDVGTENEELLKDPLYIGLRQRRVRGSEYDDFL DEFMEAVSSKYGMNCLIQFEDFANVNAFRLLNKYRNQYCTFNDDIQGTASVAVAGLLAALRITKNKLSDQTILFQ GAGEAALGIAHLIVMALEKEGLPKEKAIKKIWLVDSKGLIVKGRASLTQEKEKFAHEHEEMKNLEAIVQEIKPTA LIGVAAIGGAFSEQILKDMAAFNERPIIFALSNPTSKAECSAEQCYKITKGRAIFASGSPFDPVTLPNGQTLYPG QGNNSYVFPGVALGVVACGLRQITDNIFLTTAEVIAQQVSDKHLEEGRLYPPLNTIRDVSLKIAEKIVKDAYQEK TATVYPEPQNKEAFVRSQMYSTDYDQILPDCYSWPEEVQKIQTKVDQ

932/2825 FIGURE 821

933/2825 FIGURE 822

MASPHQEPKPGDLIEIFRLGYEHWALYIGDGYVIHLAPPSEYPGAGSSSVFSVLSNSAEVKRGRLEDVVGGCCYR VNNSLDHEYQPRPVEVIISSAKEMVGQKMKYSIVSRNCEHFVAQLRYGKSRCKQVEKAKVEVGVATALGILVVAG CSFAIRRYQKKATA

934/2825 FIGURE **823A**

GGAGGCCGGCCTAGAAACCCTCCCTCCCAGAAGAAAGCCGATCCCAGTTCAGGTGGGGTCTTCCTCGGTTGCGTA CCTGGCTGGAGCCGAGCTGGTGGCCGCCGGCAGCCGGCGTTTCTGGTGATGACAGCCCCGAAATGAAAGCAGCC CGGCCGCCGCCTCCGAGGGCTGCAGGGAGATCAGCGTCCAGCAAATAAGAAGCAAGTCCTGGACCCGGAGGAGGA CGCAGAGCTGCCTCTGGGCGCCTGACCCCGCGCTGACATCACAACCTGTGACAGGCGCATCACGCCCGGTACCTG $\tt CTCCCGGCCGCTCCTCCCAGCCTCTTTGTATGCCGCAGACAT{\tt C}GCCAGCCAGCAGGATTCGGGCTTCTT$ TGAGATCAGTATCAAATATTTACTGAAATCCTGGAGTAATACTTCTCCCGTTGGCAACGGTTACATCAAGCCTCC GGTTCCACCTGCTTCTGGCACGCACAGGGAGAAAGGGCCGCCAACCATGCTACCCATCAATGTGGACCCAGACAG TAAACCAGGAGAATATGTCCTCAAAAGTTTATTTGTCAACTTCACCACTCAGGCTGAACGCAAGATTCGTATCAT TATGGCAGAGCCCCTGGAAAAGCCATTGACGAAATCTCTGCAACGTGGAGAAGACCCCCAATTTGATCAGGTCAT CAGCTCAATGAGCTCCCTTTCTGAGTACTGCCTGCCTTCCATTCTACGTACATTATTTGACTGGTATAAAAAGGCA AAATGGCATTGAGGATGAATCACATGAATACAGACCAAGAACAAGCAATAAATCAAAAAGCGATGAACAACAGCG GATTCCACTTCATCCTGTAATAGACAGTTTAATACATGATGTTATTAACTTGGCTTTCAAGCACTTTAAATACAA AGAAGGGTACCTTGGTCCCAACACTGGCAATATGCATATTGTGGCAGACCTGTATGCAGAAGTCATTGGAGTGTT GGCACAAGCCAAATTCCCTGCTGTAAAGAAGAAATTTATGGCGGAGCTAAAAGAATTACGGCACAAAGAGCAGAA CCCATATGTGGTTCAAAGCATTATCAGCTTAATAATGGGCATGAAATTCTTTCGAATTAAGATGTATCCAGTGGA GGATTTTGAGGCCTCTCTCAGTTTATGCAGGAATGTGCACATTACTTCCTCGAGGTCAAAGACAAAGATATCAA GCATGCCTTGGCTGGGCTTTTTGTTGAAATACTTGTTCCAGTTGCTGCTGCTGTTAAAAATGAAGTAAATGTTCC CTGCCTTAGAAATTTTGTGGAAAGCCTGTATGACACCACGCTGGAACTTTCTTCTCGAAAGAAGCATTCCTTGGC ${\tt CAACAACTGCTTGTCCAACCTTAAAAACAAAGATCCCAAGATGGCTCGAGTTGCACTGGAATCTCTCTACAGATT}$ AACACTTTTCCCCAAAGGGTCCCGCGGTGTGGTACCAAGGGACATGCCTCTGAACATCTTTGTGAAAATCATCCA AAAAGCATTCAGTCTCAACCCAGAGAGAATGAACATTGGTTTACGGGCATTCTTGGTCATAGCTGATAGCTTGCA AACATATTTGAGTAAAACACTAACTGAAGAGGAAGCCAAAATGATAGGCATGTCCTTATATTACTCTCAAGTACG AAAAGCTGTAGACAACATTTTAAGGCACCTTGATAAAGAAGTAGGAAGGTGTATGATGCTGACTAATGTACAGAT GTTAAACAAAGAACCGGAAGACATGATCACGGGTGAGAAAAGCCAAAAATAGATCTTTTCAGGACCTGTGTTGC TGCTATTCCTCGACTGCTTCCTGATGGGATGTCAAAACTTGAACTTATTGACTTACTGGCTAGGCTCTCTATTCA TATGGATGATGAATTGCGACATATTGCACAAAATTCTCTTCAGGGTTTACTTGTTGACTTCTCAGATTGGAGGGA AGATGTACTATTTGGCTTTACCAACTTCCTGCTCCGGGAAGTAAATGATATGCATCACACACTCCTTGATTCGTC CCTGAAGTTGCTGCTGCAGCTGCTCACCCAGTGGAAACTAGTCATCCAGACAAGGAAAAAGTCTATGAACAAGC CAACAAAATCAGAAATTCAGAGCTCATCGCAAATGGCTCCAGTCACAGAATTCAGTCGGAACGAGGTCCCCACTG CAGTGTACTCCACGCTGTAGAAGGTTTTGCTCTGGTTTTACTCTGCAGTTTCCAGGTGGCCACACGCAAACTGTC CGTTTTAATACTCAAGGAAATTCGAGCGTTGTTTATTGCCCTGGGGCAGCCTGAGGATGACGACAGGCCGATGAT TGATGTCATGGATCAGCTAAGTTCTTCCATTCTAGAAAGTTTTATTCATGTAGCAGTTTCGGATTCAGCAACATT ACCACTCACCCACAATGTGGATCTGCAGTGGTTGGTGGAATGGAACGCAGTCCTGGTCAATAGCCATTATGATGT GAAAAGCCCTTCCCATGTCTGGATATTTGCACAGTCTGTCAAAGACCCCTGGGTCCTCTGCCTCTTCAGCTTCCT CCGGCAGGAGAACTTACCCAAGCACTGCCCCACAGCCCTCAGCTATGCCTGGCCTTATGCCTCACTCGGCTCCA GTCGGTGATGCCTCTGGTGGACCCAAATAGCCCAATTAATGCCAAGAAAACCAGCACTGCCGGCAGCGGAGACAA CTATGTTACTTTGTGGAGAAATTACCTAATTCTTTGTTTTGGAGTTGCAAAACCCAGTATTATGAGCCCAGGACA CTTAAGAGCTTCCACTCCAGAAATAATGGCGACCACACCTGATGGTACAGTGAGCTACGATAACAAGGCCATAGG CACCCCATCGGTGGGAGTTCTGTTAAAGCAGTTGGTGCCTTTGATGAGACTAGAGAGCATTGAGATCACAGAGTC CTTAGTTTTAGGATTTGGAAGAACAAATTCCCTTGTTTTCAGAGAATTGGTAGAAGAACTTCATCCATTAATGAA AGAAGCTCTGGAAAGAAGACCAGAGAACAAGAAACGCCGAGAACGGCGAGACTTGTTAAGGCTACAACTACTTCG

935/2825 FIGURE 823B

AATTTTTGAACTTTTGGCTGATGCTGGTGTAATAAGTGACACACAAATGGAGCCCTAGAGCGGGATACTTTAGC CCTGGGAGCTTTGTTCTTAGAATATGTGGACTTGACCCGCATGCTCCTAGAAGCTGAAAATGACAAAGAAGTTGA AATTCTTAAAGATATCCGGGCACATTTTAGTGCAATGGTGGCCAACTTGATTCAGTGTGTTCCAGTTCACCACCG AAGATTTCTCTTCCCCCAGCAAAGTCTGAGGCACCACCTTTTCATCTTATTCAGCCAGTGGGCAGGACCCTTCAG CATTATGTTCACTCCTCTGGATCGTTACAGTGACAGAAATCATCAGATTACAAGATATCAGTATTGTGCATTAAA AGCAATGTCAGCAGTACTGTGCTGTGGCCCTGTCTTTGACAATGTGGGCCTTTCCCCAGATGGCTACCTATATAA ATGGCTTGACAACATTCTGGCTTGTCAAGATTTACGAGTTCATCAACTTGGCTGCGAAGTTGTTGTCTTGCTACT GGAACTTAATCCTGACCAAATAAATCTTTTTAACTGGGCAATTGACCGATGCTACACAGGTTCCTACCAACTTGC ATCTGGCTGCTTCAAAGCCATAGCAACTGTGTGTGGAAGCAGGAACTATCCCTTCGACATAGTGACATTGTTAAA CCTTGTTCTATTCAAGGCCTCTGACACCAACAGAGAGATTTATGAAATCTCCATGCAGCTCATGCAGATCCTTGA AGCAAAGCTTTTTGTATACTCAAAGAAGTCGCTGAGCAAAGACCGGGAAGTATTCTCTATGGAACACACGGCCC GCTGCCACCCCTCTACAGCGTGTCACTCGCCCTCTTGTCATGTGAGCTGGCCAGGATGTACCCTGAGCTCACACT CCCCCTCTTCTCAGGTAAGCCAGCGATTCCCCACAACACACCCCAACGGGCGCCAGATCATGCTTACCTGC TGCCCTGGCTGCACAACATCGAGCTGGTGGACAGCAGCCTCCTCCTCCCGGGTCGAGCCCCAGCAGCCCAGAGGA CGAAGTCAAGGACCGGGAAGGTGACGTGACTGCTTCTCACGGGCTGAGAGGAAATGGCTGGGGCTCTCCAGAAGC CACGTCACTGGTCCTGAACAACCTCATGTACATGACGGCCAAGTATGGAGATGAAGTTCCTGGGCCAGAAATGGA AAATGCTTGGAATGCTTTAGCCAACAATGAGAAATGGAGCAACAACCTGAGGATCACCTTGCAGTTCCTGATTAG CCTCTGTGGGGTCAGCAGCGACACAGTTCTCCTACCCTATATTAAAAAAGTGGCAATATACTTGTGCCGTAACAA CACCATTCAAACCATGGAAGAGCTTCTCTTTGAGCTGCAGCAGAACCCGTGAACCCCATCGTCCAGCATTG TGACAACCCGCCTTCTACCGCTTCACGGCCAGTAGCAAGGCTTCCGCAGCAGCCTCAGGAACCACCTCTAGCAG CAATACAGTGGTTGCTGGCCAGGAAAATTTCCCAGATGCTGAGGAGAACAAGATATTGAAAGAATCTGATGAAAG GTTTAGTAATGTCATCAGAGCCCACACTCGCCTCGAGTCAAGATACAGCAATAGCTCTGGAGGATCCTACGATGA AGATAAAAATGATCCAATTTCTCCCTACACGGGCTGGTTGCTGACTATTACAGAGACCAAGCAGCCGCAGCCCTT ACCGATGCCTTGTACTGGAGGATGCTGGGCCCCCTGGTTGACTATCTCCCGGAGACCATCACTCCCCGGGGGCC ACTCCACAGGTGCAATATTGCTGTAATTTTTATGACTGAAATGGTGGTGGATCACAGTGTACGAGAAGACTGGGC GCTTCATCTACCATTATTACTTCATGCTGTCTTCTTAGGTTTAGACCACTACCGGCCTGAAGTCTTTGAACACAG CAAAAAACTGCTTCTTCACCTCTTGATTGCCCTCTCTTGCAACAGCAATTTCCATTCCATTGCTTCCGTGCTCCT TGGCTTTGACTTCCTGAGAGAGGACCAGTCATCCCCGGTGCCTGACTCAGGGCTTAGTTCAAGCTCCACCTCCTC TAGCATCAGTCTGGGAGGCAGCAGTGGAAACCTCCCACAGATGACCCAGGAGGTAGAAGATGTGGACACAGCTGC TGAAACAGATGAGAAGGCAAACAAGCTCATTGAGTTTCTCACGACCAGGGCATTTGGTCCACTTTGGTGCCATGA AGACATCACACCTAAAAATCAAAATTCAAAGAGTGCTGAACAGCTCACTAATTTTCTACGTCACGTTGTATCTGT ATTTAAAGATTCCAAATCAGGCTTCCATCTGGAGCACCAGTTGAGTGAAGTTGCATTGCAGACAGCCCTCGCAAG CTCTTCAAGGCACTATGCTGGTCGGTCCTTCCAGATATTCCGGGCCCTCAAGCAACCTCTGTCAGCACATGCCTT ATCTGACCTTCTCCAAGATTGGTGGAGGTGATAGGAGAACATGGAGATGAGATTCAGGGTTATGTAATGGAAGC GCTCCTAACCTTGGAGGCGGCTGTGGATAACTTGTCTGACTGCTTGAAGAACAGTGACCTCCTAACTGTATTGTC GAACCCGGGAACCACCAGCGCAACACCGCAACTGCCGAACGGAGCCGGCATCAACGAAGCTTCTCTGTGCCCAA GAAGTTTGGTGTCATCGACCGATCCTCTGACCCACCTCGAAGTGCCACACTGGACAGAATTCAGGCTTGTACCCA ACAAGGCCTCTCCTCAAAAACCAGAAGCTCATCCTCCTTGAAGGACAGTCTCACGGACCCATCCCACATAAACCA TCCCACCAACCTGCTGGCCACCATATTCTGGGTCACAGTGGCCTTGATGGAGTCTGATTTTGAGTTTGAATACTT AATGGCCTTAAGGCTGTTGAGCAGACTACTGGCACATATGCCACTCGATAAGGCTGAGAACCGAGAAAAGCTTGA GAAACTCCAGGCACAGCTGAAGTGGGCCGACTTCTCCGGGCTGCAGCAGCTGCTGCTGAAAGGATTCACATCCCT CACCACCAGACCTGACCCTGCAGCTCTTCAGTCTGCTGACACCAGTGTCCAAAATATCCATGGTGGATGCATC CCACGCTATTGGGTTTCCACTGAATGTCTTGTGTCTCCTGCCTCAGCTGATTCAGCATTTTGAAAATCCCAATCA ACATGTCATGACTCTTTATAAAACGCACAGCTACACGAGGGACTGTGCCACGTGGGTCAATGTGGTCTGTCGATA

936/2825 FIGURE 823C

TAGTGTGCAGCAGCCCCTGCTCCAGGTGATCTACAGTCTTCTCAGCTACATGGACCTTTCTGTCGTTCCTGTCAA ACAGTTCAATGTGGAAGTTCTGAAGACAATTGAAAAATATGTGCAAAGTGTTCACTGGAGAGAAGCTCTGAATAT CTTGAAGCTGGTAGTTTCTCGGTCAGCCAGCCTTGTTTTACCTTCATACCAGCACAGTGACCTCTCAAAAATAGA AATACATCGAGTGTGGACTAGTGCTTCCAAGGAATTACCTGGGAAAACCCTGGACTTCCACTTCGATATTTCGGA GACTCCAATCATCGGGAGGCGGTATGATGAGCTGCAGAATTCTTCTGGGCGTGATGGGAAGCCCAGGGCCATGGC CGTCACCCGGAGCACATCTTCCACTTCCTCAGGCTCCAACTCCAACGTCCTTGTTCCAGTGAGCTGGAAAAGGCC CCAGTATTCTCAGAAGAGAACAAAAGAGAAGTTGGTACATGTCCTTTCTCTGTGTGGCCAAGAAGTAGGATTGAG CAAAAATCCATCAGTGATTTTTTCATCGTGTGGGGATCTGGATCTGCTTGAGCACCAGACAAGCTTGGTATCTTC TGAGGACGGTGCCCGAGAGCAGGAGAACATGGATGACACAAACAGCGAGCAGCTTTAGAGTCTTCAGAGACTT CGACTTCCTAGATGTGGAGCTGGAGGATGGAGAGGGTGAGAGTATGGACAATTTCAACTGGGGAGTGCGCAGACG TTCTCTGGACAGCCTGGATAAGTGTGATATGCAGATTCTGGAGGAGCGCCAACTGTCAGGAAGCACTCCTAGCCT CTCAGACCTAATCATGACTCTCCCCCCTCTGAAGAGACGAATCCCATGGAGCTGCTCACCACAGCCTGTGACTC GACCCCTGCAGAACCTCATTCCTTTAACACCAGAATGTCCAGCTTTGATGCTTCCTTGCCTGATATGAATAATCT GCAGATTTCTGAGGGTTCAAAGGCTGAAGCTGTTCGTGAGGAGGACACCACCGTGCATGAGGATGATCTTTC TAGTTCCATCAATGAACTCCCAGCAGCTTTTGAATGCAGCGACAGCTTTAGCCTGGACATGACTGAGGGGGGAAGA AAAAGGCAATCGGGCACTGGACCAGTTTACCCTGGCGAGCTTTGGAGAAGGTGACAGGGGAGTCTCTCCCCCTCC CTCGCCCTTCTCCAGCCATCCTTGCCGCCTTTCAGCCCGCAGCCTGTGACGATGCCGAGGAGGCCTGGCGCAG CCACATCAACCAGCTTATGTGTGACTCAGATGGCTCCTGTGCTGTGTATACATTTCATGTGTTCTCCTCCTTGTT TAAGAATATTCAGAAAAGGTTCTGCTTCCTAACCTGTGATGCAGCCAGTTACCTTGGAGATAACCTCCGGGGAAT CGGATCCAAATTTGTCAGCTCTTCCCAGATGCTCACCTCCTGCTCTGAATGTCCTACACTTTTTGTGGATGCCGA GACTCTCCTTTCATGTGGACTTCTGGACAAGCTCAAGTTCAGTGTTTAGAACTGCAAGAATATTTTGGATACCTA CAACAACAGGAAAGAGGCCACACTCTCTTGGCTTGCAAATTGTAAGGCAACATTTGCAGGGGGGATCAAGAGATGG AGTAATTACCTGTCAACCAGGGGACTCCGAAGAAAAGCAATTGGAACTGTGTCAGAGATTATATAAGCTACACTT TCTCTATTCAGACGGCGCGTGGTCCGAGCCCACCTTCACGTCCACTGAAGCAGCCATCCAGTCCATGCTGGAGTG CCTGAAGAACAACGAACTCGGCAAAGCTTTGCGGCAGATCAGGGAGTGCAGAAGTCTGTGGCCCAATGACATCTT TACTTATGCCCTGGTGGGGTCTAACCAGAGCCTGACCGAGATCTGCACCAAGCTGATGGAGCTGAACATGGAGAT $\tt CCGGGACATGATCCGCAGGGCCCAGAGTTACCGAGTCCTCACTACTTTTCTTCCAGACTCCAGTGTTTCTGGCAC$ TAGTCTCTGACAGGAGCCTCCTGTCCCCACTGGGTTCCAAACTGCAGTGCTGCCATGCTGGGGCAACGTCATTCA GTGTCTTCTCGGCCTTCAAAAGGCTTGGACAGACTGTTCTCCCTCTTGTTACCTGTAGGGCTTTTTCTAAAGAGG ATGGCAGAACTTCCAACGTGTAGCAATACTATAAGAACCAAGGTAGCTTAGAACGTCCTGGACAGACTCCACTCA TCATGCTGTGTGGCACAAATGTGTTACATTTGACCGAGCATATGCAACTCGCTACTGAAGAAGTGACTTCCGTTG CATACCAAAGCCGACTACACTGAACAGTACCTTCCTTTCTAGAAACAATTTTAGATTGGCAAAAGTGCAATGTTT TCTTCACTCAAAAAATTTTATATTCTCAAACATGTATATTCTTTCCCTGTCTTGTTCCATTTTCTTTTTTT ATGGTCAAAAGCTGACAGCCTGTGTATGTGAAAAGGGAATTGTAAATGGACTGCAATGTAATGTACACTGTAATT TGAATACAATTACTGTATCTAAAAGGAGCTGCTATGAAGTACCTTTCTTATGTTGCTAGGCTACTGTTTCTGAAA GCCCTGGATCTCTTTGCACCAAAAATGGTCCAGATAGACTCTTTTTAAGGATCTTGGCTGCTTTTTACTAGAAGG AGTAGAATCATAAAGGTAAAATATGAGTTGTTACTTTGTTTCTTCGATGTCATATTTTATGTGTAATATATGT AAAGGGCCATTCTTAAGTTCTCCCTTAAACTTAATGCTGTCAAGTGTTAGATGTGTGCATGTGAACTTGTTGCA CTGCAGAAACATATTCAGAGTTTATCTATGTAACTTATTCACTCTGTAAATACATTTAAAGTTTTTGTGATGTAA

937/2825 FIGURE **824**

MASQQDSGFFEISIKYLLKSWSNTSPVGNGYIKPPVPPASGTHREKGPPTMLPINVDPDSKPGEYVLKSLFVNFT TQAERKIRIIMAEPLEKPLTKSLQRGEDPQFDQVISSMSSLSEYCLPSILRTLFDWYKRQNGIEDESHEYRPRTS NKSKSDEQQRDYLMERRDLAIDFIFSLVLIEVLKQIPLHPVIDSLIHDVINLAFKHFKYKEGYLGPNTGNMHIVA DLYAEVIGVLAQAKFPAVKKKFMAELKELRHKEQNPYVVQSIISLIMGMKFFRIKMYPVEDFEASLQFMQECAHY FLEVKDKDIKHALAGLFVEILVPVAAAVKNEVNVPCLRNFVESLYDTTLELSSRKKHSLALYPLVTCLLCVSQKQ LFLNRWHIFLNNCLSNLKNKDPKMARVALESLYRLLWVYMIRIKCESNTATQSRLITIITTLFPKGSRGVVPRDM PLNIFVKIIQFIAQERLDFAMKEIIFDFLCVGKPAKAFSLNPERMNIGLRAFLVIADSLQQKDGEPPMPVTGAVL PSGNTLRVKKTYLSKTLTEEEAKMIGMSLYYSQVRKAVDNILRHLDKEVGRCMMLTNVQMLNKEPEDMITGERKP KIDLFRTCVAAIPRLLPDGMSKLELIDLLARLSIHMDDELRHIAQNSLQGLLVDFSDWREDVLFGFTNFLLREVN DMHHTLLDSSLKLLLQLLTQWKLVIQTQGKVYEQANKIRNSELIANGSSHRIQSERGPHCSVLHAVEGFALVLLC SFQVATRKLSVLILKEIRALFIALGQPEDDDRPMIDVMDQLSSSILESFIHVAVSDSATLPLTHNVDLQWLVEWN AVLVNSHYDVKSPSHVWIFAQSVKDPWVLCLFSFLRQENLPKHCPTALSYAWPYAFTRLQSVMPLVDPNSPINAK KTSTAGSGDNYVTLWRNYLILCFGVAKPSIMSPGHLRASTPEIMATTPDGTVSYDNKAIGTPSVGVLLKQLVPLM RLESIEITESLVLGFGRTNSLVFRELVEELHPLMKEALERRPENKKRRERRDLLRLQLLRIFELLADAGVISDST NGALERDTLALGALFLEYVDLTRMLLEAENDKEVEILKDIRAHFSAMVANLIQCVPVHHRRFLFPQQSLRHHLFI LFSQWAGPFSIMFTPLDRYSDRNHQITRYQYCALKAMSAVLCCGPVFDNVGLSPDGYLYKWLDNILACQDLRVHQ LGCEVVVLLLELNPDQINLFNWAIDRCYTGSYQLASGCFKAIATVCGSRNYPFDIVTLLNLVLFKASDTNREIYE ISMQLMQILEAKLFVYSKKVAEQRPGSILYGTHGPLPPLYSVSLALLSCELARMYPELTLPLFSGKPAIPHNTPQ RAPDHAYLPAALAAQHRAGGQQAPPPGSSPSSPEDEVKDREGDVTASHGLRGNGWGSPEATSLVLNNLMYMTAKY GDEVPGPEMENAWNALANNEKWSNNLRITLOFLISLCGVSSDTVLLPYIKKVAIYLCRNNTIQTMEELLFELQOT EPVNPIVQHCDNPPFYRFTASSKASAAASGTTSSSNTVVAGQENFPDAEENKILKESDERFSNVIRAHTRLESRY SNSSGSYDEDKNDPISPYTGWLLTITETKQPQPLPMPCTGGCWAPLVDYLPETITPRGPLHRCNIAVIFMTEMV VDHSVREDWALHLPLLLHAVFLGLDHYRPEVFEHSKKLLLHLLIALSCNSNFHSIASVLLQTREMGEAKTLTVQP AYOPEYLYTGGFDFLREDOSSPVPDSGLSSSSTSSSISLGGSSGNLPOMTOEVEDVDTAAETDEKANKLIEFLTT RAFGPLWCHEDITPKNQNSKSAEQLTNFLRHVVSVFKDSKSGFHLEHQLSEVALQTALASSSRHYAGRSFQIFRA LKQPLSAHALSDLLSRLVEVIGEHGDEIQGYVMEALLTLEAAVDNLSDCLKNSDLLTVLSRSSSPDLSSSSKLTA SRKSTGQLNMNPGTTSGNTATAERSRHQRSFSVPKKFGVIDRSSDPPRSA

938/2825 FIGURE 825

GACGGCGAGACCCCGCCCCATCCCCGACTGCCTGAACCGCGCCAGGAGACCGCAAGTCCAGCGTACCCACA $\verb|CCGACGGTGGTGTCAGCAAAAGCACAACATCTGGGGAGGAGCTAGTAGTCCAGGTTCCCGTAGTGGATGTGCAAA| \\$ GCAACAACTTCAAGGAGATGTGGCCATCCCTCCTGCTAGCCATAAAGACAGCTAATTTCGTGGCTGTGGACACGG AGCTGAGTGGGCTTGGGGACAGGAAGAGTTTGCTGAACCAGTGCATTGAGGAACGTTACAAGGCCGTGTGTCATG CTGCCAGGACCCGTTCTATCCTTTCCCTGGGCCTCGCCTGCTTCAAGCGGCAGCCAGACAAGGGTGAACATTCCT ATCTGGCTCAAGTGTTCAATCTCACTCTGCTGTGCATGGAGGAGTATGTCATAGAACCAAAGTCTGTGCAGTTCC TGATACAGCATGGCTTCAACTTCAACCAGCAGTATGCCCAAGGCATCCCCTACCATAAGGGCAATGACAAGGGTG ATGGCCTTATAGACTTGGTGTTCCTGTACCAGAACTTCTATGCACACCTCCCTGAGAGTCTGGGAACCTTCACCG CTGACCTGTGTGAGATGTTCCCAGCAGGCATTTATGACACCAAATATGCTGCTGAGTTTCGTGCCCGTTTCGTGG CCTCCTACTTAGAATATGCCTTCCGGAAATGTGAACGGGAAAATGGGAAGCAGCGGGCAGCTGGCAGCCCACACC CCCACCGTCCTCATCCCACCAGCATCTGTGACAACTTCTCGGCTTATGGCTGGTGCCCCCTGGGACCACAGTGTC CTCAGTCTCACGATATTGACCTTATCATTGACACTGATGAGGCTGCGGCAGAGGACAAGCGGCGACGGCGACGAC GTAGGGAAAACGGAAGAGGGCTTTATTGAACCTACCGGGGACACAGACCTCTGGGGAAGCTAAGGATGGTCCTC CCAAGAAGCAGGTCTGTGGGGATAGCATCAAGCCTGAAGAAACCGAGCAGGAGGTGGCTGCCGATGAAACTAGGA ACCTGCCTCACTCCAAGCAAGGCAACAAAAATGACTTAGAGATGGGGGATTAAGGCAGCAAGGCCTGAAATAGCTG GTTTTGATGCCTTTATGACAGGTTATGTGATGGCCTATGTGGAAGTGAGCCAGGGACCGCAGCCCTGCAGCTCTG GACCCTGGCTCCCTGAATGCCACAATAAGGTATATTTGAGTGGCAAAGCTGTACCCCTCACAGTGGCCAAGAGCC AGTTCTCTCGTTCCTCCAAAGCCCACAATCAGAAGATGAAGCTCACTTGGGGCAGTAGC**TGA**TGCAACTTCCACC TTGCTCTCAGGTGGAACAGAGGTATTTTGGGTCTCTCTAGCCTGAAATGTCATCCTCAACTGCTACTGAGTTTAG GGGAGGGGAATGTCTTGACAGACATCACTGCATTGCCCTGGACCGCCTCCTTTATCCCAGTGTTTGAGGTACAA GTAAGAAGGCTGACCAGCACCTGTAACACTGACTTTATTTTTAAGTCTGAAAATGTCTTGGGAAAGTTTTAC

939/2825 FIGURE 826

MAADSDDGAVSAPAASDGGVSKSTTSGEELVVQVPVVDVQSNNFKEMWPSLLLAIKTANFVAVDTELSGLGDRKS LLNQCIEERYKAVCHAARTRSILSLGLACFKRQPDKGEHSYLAQVFNLTLLCMEEYVIEPKSVQFLIQHGFNFNQ QYAQGIPYHKGNDKGDESQSQSVRTLFLELIRARRPLVLHNGLIDLVFLYQNFYAHLPESLGTFTADLCEMFPAG IYDTKYAAEFRARFVASYLEYAFRKCERENGKQRAAGSPHLTLEFCNYPSSMRDHIDYRCCLPPATHRPHPTSIC DNFSAYGWCPLGPQCPQSHDIDLIIDTDEAAAEDKRRRRRREKRKRALLNLPGTQTSGEAKDGPPKKQVCGDSI KPEETEQEVAADETRNLPHSKQGNKNDLEMGIKAARPEIADRATSEVPGSQASPNPVPGDGLHRAGFDAFMTGYV MAYVEVSQGPQPCSSGPWLPECHNKVYLSGKAVPLTVAKSQFSRSSKAHNQKMKLTWGSS

940/2825 FIGURE 827A

TTTTTTTTTTTTTTACAGCTGTTCCTGCTGATTGCTTTTTAAGCTAGGCTCAAGTGTGGGGCCACGGAATCCTT CAGGTGCCCTTCAAATCAGACATTACATTTCAGCTCTCAGGGCTAGAATGCTTCCCCCTCTTATCTTGGGCAGGT CTTTGAAGAAGCCAGTTTAGCCTTGGTAATTTTGAATTTTGGCCATAGTATACGGAGGAGAGCTAGGTAGCTATCA GTGGGCAAAGGAGTGTGTTTTGGGGCAGGGTACTGAGTCCCATATGTAATCAAGAGACCTGGAGGGCCCAAGGA AGTCAGTGACACCCCCACAAGCTACATATCCTCCAATGGATTGTGGTGGTCATCTCCGGCTACCGAAGCCACCTT GCTTATTCCCATAGGCTTACAAACATTATGATGCTTCACACGAGTTGTGGGTGAAGTGAAGGAATTGGAAAGTCC CGGTCTAAGGCCCTGCAAAACTGCTGTCTGGTAAAAACAAGTGACTATGGCAAGGCCCTGAATGTACTACGTATA GACTTAAGTCAGTATTTGTGCCTAACAGCAAAGCATGCCACAAGAGGCTCCTCAAAAGCCGGGTAATTGATGCCA TTGTTTTCTTGACCTAGTTGAAGAGTCAAAGAAGAAGATGATGCAGAAACTTCTTTCCAGTGAAGTGACAGAT GGACCACATATTGTCGGTGTTAACCACTGTTATCACTGGCCCAGGTCGAGTGAGGACGAGTCATTTCCCCACTAA GCACTTGGGGAGTGGCAGGGACCACATATTGTCAGTGTTAACCACTGTTATGACTGGCCCAGGTCGGGTGACGAC GAGCCATTTCCGCACTATTGTCCCTTCTTCCTCACTGCACCCCCAGCCCTGCACCCTCAGCTCCAAATACACATT CCCGGTGACAGATTTTTGGTTTTGGTGGTGGCTTTTTGGGCAAGGACAGGTCAGTTCATAAAAGATGACGCCCCAG GTCTGGATCGAATTCTCCTGGGAGCCAGACCAAAGCACCTTCCTGGACTGCAAAGTGTGGAAGCCGCCGCGGGCC GGGCGCGGGGAGGTGTC**ATG**CGCCGGAACCTGCGCTTGGGGCCAAGCTCTGGAGCTGACGCGCAGGGGCAAGGCG CCCCGCGTCCCGGACTGCCGCGCTCCCCACCGCGTCGCAGGCCTCGAGAGGCTCCGGAAGTA CTGGGTGCAGCCTGATGGCGCAGGAGGTAGACACGGCACAGGGCGCCGAGATGCGGCGGGGCGCGGGCGCGCTC GCATCCCCTCCCGGCGCCACTGGCCGGTGCCCTACAAGCGCTTTGACTTCCGTCCAAAACCTGATCCTTATTGTC ${\tt AAGCTAAGTATACTTTCTGTCCAACTGGCTCACCTATCCCAGTTATGGAGGGTGATGATGACATTGAAGTTTTTC}$ GATTACAAGCCCCAGTATGGGAATTTAAATATGGAGACCTCCTGGGACACTTGAAAATTATGCATGATGCCATTG GATTCAGAAGTACATTAACTGGCAAGAACTACACAATGGAATGGTATGAACTTTTCCAACTTGGCAACTGTACAT ATGATGTTCACTGGAAGGAAAATGGGACATTAGTTCAAGTAGCAACTATATCAGGAAACATGTTCAACCAAATGG GGGCAGAGACATGGTTTGATTCCTACGACTGTTCCAAATTTGTGTTAAGGACCTTTAACAAGTTGGCTGAATTTG GAGCAGAGTTCAAGAACATAGAAACCAACTATACAAGAATATTTCTTTACAGTGGAGAACCTACTTATCTGGGAA ATGAAACATCTGTTTTTGGGCCAACAGGAAACAAGACTCTTGGTTTAGCCATAAAAAGATTTTATTACCCCTTCA AACCACATTTGCCAACTAAAGAATTTCTGTTGAGTCTCTTGCAAATTTTTGATGCAGTGATTGTGCACAAACAGT TCTATTTGTTTTATAATTTTGAATATTGGTTTTTACCTATGAAATTCCCTTTTATTAAAATAACATATGAAGAAA ${\tt TCCCTTTACCTATCAGAAACAAAACACTCTCTGGTTTA}{\tt TAAA}{\tt AACACCTTAATTCTACTGCTCTTTTTTCTCCAAT$ ${\tt CACCAGCATCTGTTTTCAGGGGGTGATTTTACTTTTGTGAATTCCTTAGCCTTTCTTCCTTGGTGCATAAAGTT}$ AAAATGCACATCAGCAGAATTGCTGCATATTAACATCTCAGGACTCTTCTCTTGTAAAGAAGCTGAAATTCGTAC TATATTGGCCAAAGTGAGCGAGTTAGGTGATCTTGGTTTCAATTTCCGAGCCTTTGTTAATATGGAGAATTATGG TTĆACTTAAGTGCTTTAGTTTAGACTAACTTTACAACTTCTATAACTTTTGGAACCAAGTTTAGTATAGTCTGAT TACATTCCATTCACCTAACTTTAGACATTCGTTTAGACACCATAACTGGAGTGATTGTGCTTCTAGATGTGGCAA AACATCACTAAATCTTAGGATTTAGGATTGCTGTAAAGATGTTAAGTTGTGTATGTTTTGGCAGGTCACATTGAATG GCAGTGATAATGATTAATCAAAGAACAAATGTCATCCTTGATCTTGCCTAATGTAGTTTATGTGCCAAACTTTCC AGGGTTTTGTAGTCACCTAGATTTTAAGCTGATAGCATAGTGCTTCAGCGGTTCTTCTAACCGGGGTATGCAGGA ACATGGCTGCAGACACGTTTGGGTAAACAGGCACCTTCTGACTTCTTCATTGTTTCCTGTAGTTCTCCCTCTTCC GCTGTGCTTTAATTAGAGCCAGACAAGCTTTCAAGGTCCTTTAAGTATTTGATGATCAACTGAACACGTTTCTAT

941/2825 FIGURE 827B

942/2825 FIGURE 828

MRRNLRLGPSSGADAQGQGAPRPGLAAPRMLLPPASQASRGSGSTGCSLMAQEVDTAQGAEMRRGAGAARGRASW CWALALLWLAVVPGWSRVSGIPSRRHWPVPYKRFDFRPKPDPYCQAKYTFCPTGSPIPVMEGDDDIEVFRLQAPV WEFKYGDLLGHLKIMHDAIGFRSTLTGKNYTMEWYELFQLGNCTFPHLRPEMDAPFWCNQGAACFFEGIDDVHWK ENGTLVQVATISGNMFNQMAKWVKQDNETGIYYETWNVKASPEKGAETWFDSYDCSKFVLRTFNKLAEFGAEFKN IETNYTRIFLYSGEPTYLGNETSVFGPTGNKTLGLAIKRFYYPFKPHLPTKEFLLSLLQIFDAVIVHKQFYLFYN FEYWFLPMKFPFIKITYEEIPLPIRNKTLSGL

943/2825 FIGURE 829

ACGCCGCGCATCGTGCTCATCCGCAGCCGCAAAGTGGGGCTCATGAACCGCGCCGTGCAACTGCTCATCCTGGCC TACGTCATCGGGTGGTGTTTGTGTGGGAAAAGGGCTACCAGGAAACTGACTCCGTGGTCAGCTCCGTTACGACC AAGGTCAAGGGCGTGGCTGTGACCAACACTTCTAAACTTGGATTCCGGATCTGGGATGTGGCGGATTATGTGATA $\verb|CCAGCTCAGGAGGAAAACTCCCTCTTCGTCATGACCAACGTGATCCTCACCATGAACCAGGACACAGGGCCTGTGC|\\$ CCCGAGATTCCAGATGCGACCACTGTGTGTAAATCAGATGCCAGCTGTACTGCCGGCTCTGCCGGCACCCACAGC AACGGAGTCTCAACAGGCAGGTGCGTAGCTTTCAACGGGTCCGTCAAGACGTGTGAGGTGGCGGCCTGGTGCCCG GTGGAGGATGACACACGTGCCACAACCTGCTTTTTTAAAGGCTGCAGAAAACTTCACTCTTTTGGTTAAGAAC AACATCTGGTATCCCAAATTTAATTTCAGCAAGAGGGAATATCCTTCCCAACATCACCACTACTTACCTCAAGTCG TGCATTTATGATGCTAAAACAGATCCCTTCTGCCCCATATTCCGTCTTGGCAAAATAGTGGAGAACGCAGGACAC AGTTTCCAGGACATGGCCGTGGAGGGAGGCATCATGGGCATCCAGGTCAACTGGGACTGCAACCTGGACAGAGCC GCCTCCCTCTGCTTGCCCAGGTACTCCTTCCGCCGCCTCGATACACGGGACGTTGAGCACAACGTATCTCCTGGC ATCCGCTTCGACATCATTGTGTTTTGGGAAGGCAGGGAAATTTGACATCATCCCCACTATGATCAACATCGGCTCT GGCCTGGCACTGCTAGGCATGGCGACCGTGCTGTGTGACATCATAGTCCTCTACTGCATGAAGAAAAGACTCTAC ${\tt TATCGGGAGAAGAAATATAAATATGTGGAAGATTACGAGCAGGGTCTTGCTAGTGAGCTGGACCAG{\tt TGA}_{\tt GGCCTA}}$ GTGTGCAGGATCTGTTTGCCCACTCGGCCCAGGAGGTCA

944/2825 FIGURE 830

MAGCCSALAAFLFEYDTPRIVLIRSRKVGLMNRAVQLLILAYVIGWVFVWEKGYQETDSVVSSVTTKVKGVAVTN TSKLGFRIWDVADYVIPAQEENSLFVMTNVILTMNQTQGLCPEIPDATTVCKSDASCTAGSAGTHSNGVSTGRCV AFNGSVKTCEVAAWCPVEDDTHVPQPAFLKAAENFTLLVKNNIWYPKFNFSKRNILPNITTTYLKSCIYDAKTDP FCPIFRLGKIVENAGHSFQDMAVEGGIMGIQVNWDCNLDRAASLCLPRYSFRRLDTRDVEHNVSPGYNFRFAKYY RDLAGNEQRTLIKAYGIRFDIIVFGKAGKFDIIPTMINIGSGLALLGMATVLCDIIVLYCMKKRLYYREKKYKYV EDYEQGLASELDQ

945/2825 FIGURE 831

GCCCTTGGTGCTGCAGGCGCGGTGGGCTCCGGGCCCAGGCACCGAGGGGGGCACTGGATGACTCTCCAGGTGCAGG ACCCTGCCATCTATGACTCCAGGTCTTCAGCACCCACCGTGGTACAGCGCCCCGGGATGCCGTCTGGAGCC CGGATGCCCCACCAGGGGGCCCCATGGGCCCCCGGGCTCCCCGTACATGGGCAGCCCCGCCGTGCGACCCGGC CTGGCCCCGCGGGCATGGAGCCCGCCCGCAAGCGAGCAGCGCCCCGGCCCGGGCAGAGCCAGAGCCAGAGCCAG GGCCAGCCGGTGCCCACCGCCCCCGCGGGAGCCGCAGTGCCAAGAGGAGGAAGATGGCTGACAAAATCCTCCCT CAAAGGATTCGGGAGCTGGTCCCCGAGTCCCAGGCTTACATGGACCTCTTGGCATTTGAGAGGAAACTGGATCAA ACCATCATGCGGAAGCGGGTGGACATCCAGGAGGCTCTGAAGAGCCCATGAAGCCAAAAGCGGAAGCTGCGACTC TATATCTCCAACACTTTTAACCCTGCGAAGCCTGATGCTGAGGATTCCGACGCAGCATTGCCTCCTGGGAGCTA $\tt CGGGTGGAGGGGAAGCTCCTGGATGATCCCAGCAAACAGAAGCGGAAGTTCTCTTCTTCTTCAAGAGTTTGGTC$ ATCGAGCTGGACAAGATCTTTATGGCCCTGACAACCACCTCGTTGAGTGGCATCGGACACCCACGACCCAGGAG CAGGCCCTGTGGCAGTATGTGAAGACCAACAGGCTGCAGGACTCCCATGACAAGGAATACATCAATGGGGACAAG TATTTCCAGCAGATTTTTGATTGTCCCCGGCTGAAGTTTTCTGAGATTCCCCAGCGCCTCACAGCCCTGCTATTG CCCCTGACCCAATTGTCATCAACCATGTCATCAGCGTGGACCCTTCAGACCAGAAGAAGACGGCGTGCTATGAC ATTGACGTGGAGGTGGAGGAGCCATTAAAGGGGCAGATGAGCAGCTTCCTCCTATCCACGGCCAACCAGCAGGAG CTAAGCTTCTCCAGAGACCCCAAAGGCTATGTCCAAGACCTGCTCCGCTCCCAGAGCCGGGACCTCAAGGTGATG ACAGATGTAGCCGGCAACCCTGAAGAGGAGCGCCGGGCTGAGTTCTACCACCAGCCCTGGTCCCAGGAGGCCGTC

946/2825 FIGURE 832

MTPGLQHPPTVVQRPGMPSGARMPHQGRPWAPRAPRTWAAPPCDPAWPPRDGARPQASSAPARQSQAQSQGQPEP TAPARSRSAKRRKMADKILPQRIRELVPESQAYMDLLAFERKLDQTIMRKGVDIQEALKRPMKQKRKLRLYISNT FNPAKSDAEDSDGSIASWELRVEGKLLDDPSKQKRKFSSFFKSLVIELDKDLYGPDNHLVEWHRTPTTQETDGFQ VKRPGDLSVRCTLLLMLDYQPPQFKLDPRLARLLGLHTQSRSAIVQALWQYVKTNRLQDSHDKEYINGDKYFQQI FDCPRLKFSEIPQRLTALLLPPDPIVINHVISVDPSDQKKTACYDIDVEVEEPLKGQMSSFLLSTANQQEISPLD SKIHETIESINQLKIQRDFMLSFSRDPKGYVQDLLRSQSRDLKVMTDVAGNPEEERRAEFYHQPWSQEAVSRYFY CKIQQRRQELEQSLVVRNT

947/2825 FIGURE **833**

948/2825 FIGURE 834

MCSRGWDSCLALELLLPLSLLVTSIQGHLVHMTVVSGSNVTLNISESLPENYKQLTWFYTFDQKIVEWDSRKSK YFESKFKGRVRLDPQSGALYISKVQKEDNSTYIMRVLKKTGNEQEWKIKLQVLDPVPKPVIKIEKIEDMDDNCYL KLSCVIPGESVNYTWYGDKRPFPKELQNSVLETTLMPHNYSRCYTCQVSNSVSSKNGTVCLSPPCTLARSFGVEW IASWLVVTVPTILGLLLT

949/2825 FIGURE 835

CGCCTTCCCTCCAATCAGCACCGGGGCCGGCTAGCCAGGGGCCGGCGCGCGGGGTGTGTGAGGACGCGCTCCCA GTCGCTGAGTGCCTGAGCCGGGAAGCAGTTGCTGTGGTACCTGCGCTGCCCGAGCGGAGTGGATGGCAGAGGTGG GCTGTAGAGCCAAAGTGGGGTGGGAGCGCGAAG<u>ATC</u>GCAGCTGCTGAGGAGGAGCCCGAAGCCCAAAAAGCTGAAG GTGGAGGCGCCGCAAGCGCTGAGAGAAAATATTCTCTTTTGGAATGGGAAATCCTCTGCTTGACATCTCTGCTGTA GTGGACAAAGATTTCCTTGATAAGTATTCTCTGAAACCAAATGACCAAATCTTGGCTGAAGACAAACACAAGGAA CTGTTTGATGAACTTGTGAAAAAATTCAAAGTCGAATATCATGCTGGTGGCTCTACCCAGAATTCAATTAAAGTG ATCCTGAAGAGAAAAGCTGCTGAAGCCCATGTGGATGCTCATTACTACGAGCAGAATGAGCAGCCAACAGGAACT TGTGCTGCATGCATCACTGGTGACAACAGGTCCCTCATAGCTAATCTTGCTGCTGCCAATTGTTATAAAAAGGAA AAACATCTTGATCTGGAGAAAAACTGGATGTTGGTAGAAAAAGCAAGAGTTTGTTATATAGCAGGCTTTTTTCTT CACGTTTCCCCAGAGTCAGTATTAAAGGTGGCTCACCATGCTTCTGAAAACAACAGGATTTTCACTTTGAATCTA TCTGCACCGTTTATTAGCCAGTTCTACAAGGAATCATTGATGAAAGTTATGCCTTATGTTGATATACTTTTTGGA AATGAGACAGAAGCTGCCACTTTTGCTAGAGAGCAAGGCTTTGAGACTAAAGACATTAAAGAGATAGCCAAAAAG ACACAAGCCCTGCCAAAGATGAACTCAAAGAGGCAGCGAATCGTGATCTTCACCCAAGGGAGAGATGACACTATA GGCCACTATGCAGCAAGCATCATAATTAGACGGACTGGCTGCACCTTTCCTGAGAAGCCAGACTTCCAC**TGA**TGG AAGAGCTGAAAACACAAGCCCAGGAGTGCAGACACTGCCCTAATTGCTTCCTGACAATTCCCATATTAATAAAGA AGAAAATTATCTGCCATTTTTTCCTACTATAATAATGCTGAATCTTAATTTAGAGGGTACAAGGGTATGGTAATG CTTGTAGAATCTTTATTATCTCAACAATCTAAAAAATGATGTTTATTTCCATAGTTTGATAGTGCCACTTAAATG CCAATTAAACAAGAATATAACATTTCAATAGAAATTTTTATTTCATTTCAATTACTTTGTAAATTCGTGTGTAT TTAGTACACTGATTTGTTTTTTACATTCTGCTTTGAATGCAGATGCAATTTAATAATAGATTTTTTAATGAA TTAATCTTAACATAGTAATCTTTAGCTTTTTATACAAATATATTTAATTTAGGAGTATATGTGTGTCTATACACA CACATACATAAATATACCACATATACACCTGATAGTCAAATAAGGTACAGAAATTTTATCTTGTCAATTATGCCA

950/2825 FIGURE 836

MAAAEEEPKPKKLKVEAPQALRENILFGMGNPLLDISAVVDKDFLDKYSLKPNDQILAEDKHKELFDELVKKFKV EYHAGGSTQNSIKVAQWMIQQPHKAATFFGCIGIDKFGEILKRKAAEAHVDAHYYEQNEQPTGTCAACITGDNRS LIANLAAANCYKKEKHLDLEKNWMLVEKARVCYIAGFFLHVSPESVLKVAHHASENNRIFTLNLSAPFISQFYKE SLMKVMPYVDILFGNETEAATFAREQGFETKDIKEIAKKTQALPKMNSKRQRIVIFTQGRDDTIMATESEVTAFA VLDQDQKEIIDTNGAGDAFVGGFLSQLVSDKPLTECIRAGHYAASIIIRRTGCTFPEKPDFH

951/2825 FIGURE 837

GGTGTGTCCCGGGTGTGGGGAGGCGACAGAGCCCTGGCACTTGAGGGTTGAGGGGGCCTCCCCAGCGCGCGAAC TGTGTGACCTCGGGTAAATTAGCCTTGGAACGTCAGTTTCTTCGTCTCTATAATTGAAATAATAATAGTACCTCT $\verb|CTCAGGATTGTTGTGAGCCGTCAGTGAAACACTTAGAGCAGTTTCTGGCACATGGTAGAATTGGGCTATTTGCTT|\\$ CAAATTGCTGCATGGTACTTATTTTCTCAGAAAATTCTCTAAGCCAACTTCTGCCTTGTATCCATTTTTGGGTAT CCGCTTTGCAGAGTATTCCAGTAGTCTTCAGAAACCAGTGGCTTCTCCTGGCAAAGCCTCCTCACAGAGGAAGAC TGAAGGGGATTTGCAAGGAGATCACCAGAAAGAAGTTGCTTTGGATATAACTTCTTCTGAGGAGAAGCCTGATGT TAGTTTCGATAAAGCAATTAGAGATGAAGCAATATACCATTTTAGGCTTTTGAAGGATGAAATTGTGGATCATTG GAGAGGACCGGAAGGCCACCCTCTGCATGAGGTCTTGCTGGAACAAGCCAAGGTTGTCTGGCAATTCCGGGGGAA AGAAGATTTGGATAAGTGGACAGTGACTTCTGATAAGACGATTGGAGGCAGAAGTGAAGTGTTTTTGAAAATGGG CAAGAATAACCAAAGTGCACTGCTATATGGAACTCTGAGCTCTGAGGCGCCTCAGGACGCGAGTCTTCCCGAAG TGGGTACTGTGCAATGATATCCAGGATTCCAAGGGGTGCTTTTAAGAGGGAAGATGTCTTACGATTGGTCCCAGTT CAATACTCTGTATCTCCGTGTACGTGGGGATGGTCGGCCTTGGATGGTGAATATCAAGGAGGACACAGATTTCTT CCAGAGGACGAATCAGATGTATAGTTACTTCATGTTCACCCGCGGGGGACCCTACTGGCAGGAGGTCAAGATTCC TTTTTCCAAATTTTTCTTCTCTAATCGAGGAAGAATCCGGGATGTTCAGCATGAGCTTCCGCTTGATAAGATCTC TGATCCAGCTCATACAGAAGAATTTGCCTATGAAAATTCTCCAGAGCTTAACCCAAGGCTTTTTAAA**TAA**AGATC ATATGGTAGTTTTGTTTTACTAATCTAAGGGTACTAGCATCTACAATGATATAGACAAAATAAAATATTTCTTTA ATGGCATCCAAA

952/2825 FIGURE 838

MALVHKLLHGTYFLRKFSKPTSALYPFLGIRFAEYSSSLQKPVASPGKASSQRKTEGDLQGDHQKEVALDITSSE EKPDVSFDKAIRDEAIYHFRLLKDEIVDHWRGPEGHPLHEVLLEQAKVVWQFRGKEDLDKWTVTSDKTIGGRSEV FLKMGKNNQSALLYGTLSSEAPQDGESSRSGYCAMISRIPRGAFKRKMSYDWSQFNTLYLRVRGDGRPWMVNIKE DTDFFQRTNQMYSYFMFTRGGPYWQEVKIPFSKFFFSNRGRIRDVQHELPLDKISSIGFTLADKVDGPFFLEIDF IGVFTDPAHTEEFAYENSPELNPRLFK

953/2825 FIGURE 839

GCGACGGCCCGAGCGCTTCTCCTTCGCCGAGGTTGCGCCGCCCCCTCCTCGCCGTGCCCCGGGCCCGCGCGGGATC CGGCGTCCACGTCCCCCGCTGCACCCCGCCCCGGCGCGAGAGAGCGCGAGAGCCCCAGCCGCGGGCGG GTTCGAGCCCCAGAGCCGTCCGCGATCCTGTACGTGGCCCCTGCAAAGGCCGGAGCTCCAAGCGAGCCCTGCCAA GCCCTCGGGGGACGCCGCCGACTCCATGATCCCCGAGGAGGACGATGAAGACGACGACGACGGCGGGGG ACGGGCCGGCTCGGCCATGGCGATCGGCGGCGGCGGCGGGGGCGCACGCTGGGCTCCGGGCTGCTCCTTGAGGA GCCGAGGAAATGTTCGTCGCGGCGGAACGCCTGGGGAAACCTGTCCTACGCGGACCTGATCACCCGCGCCATCGA GAGCTCCCCGGACAAACGGCTCACTCTGTCCCAGATCTACGAGTGGATGGTGCGTTGCGTGCCCTACTTCAAGGA TAAGGGCGACAGCACCAGCTCTGCCGGCTGGAAGAACTCCATCCGGCACAACCTGTCACTGCATAGTCGATTCAT GCGGGTCCAGAATGAGGGAACTGGCAAGAGCTCTTGGTGGATCATCAACCCTGATGGGGGGAAGAGGCGGAAAAGC CCCCGGCGGCGGCTGTCTCCATGGACAATAGCAACAAGTATACCAAGAGCCGTGGCCGCGCGCAGCAAGAAGAA GTCACGCAGCAGTGATGAGCTGGATGCGTGGACGGACTTCCGTTCACGCACCAATTCTAACGCCAGCACAGTCAG TGGCCGCCTGTCGCCCATCATGGCAAGCACAGAGTTGGATGAAGTCCAGGACGATGATGCGCCTCTCTCGCCCAT TATGGCAGGCACCATGAATCTGAATGATGGGCTGACTGAAAACCTCATGGACGACCTGCTGGATAACATCACGCT CCCGCCATCCCAGCCATCGCCCACTGGGGGACTCATGCAGCGGAGCTCTAGCTTCCCGTATACCACCAAGGGCTC GGGCCTGGGCTCCCCAACCAGCTCCTTTAACAGCACGGTGTTCGGACCTTCATCTCTGAACTCCCTACGCCAGTC TCCCATGCAGACCATCCAAGAGAACAAGCCAGCTACCTTCTCTCCATGTCACACTATGGTAACCAGACACTCCA GGACCTGCTCACTTCGGACTCACTTAGCCACAGCGATGTCATGATGACACAGTCGGACCCCTTGATGTCTCAGGC CAGCACCGCTGTGTCTGCCCAGAATTCCCGCCGGAACGTGATGCTTCGCAATGATCCGATGATGTCCTTTGCTGC CCAGCCTAACCAGGGAAGTTTGGTCAATCAGAACTTGCTCCACCACCAGCACCAAACCCAGGGCGCTCTTGGTGG TGCAAACCTGCCGTCATGGGCCATGAGAAGTTCCCCAGCGACTTGGACCTGGACATGTTCAATGGGAGCTTGGA ATGTGACATGGAGTCCATTATCCGTAGTGAACTCATGGATGCTGATGGGTTGGATTTTAACTTTGATTCCCTCAT CTCCACACAGAATGTTGTTGGTTTGAACGTGGGGAACTTCACTGGTGCTAAGCAGGCCTCATCTCAGAGCTGGGT GCCAGGCTGAAGGATCACTGAGGAAGGGGAAGTGGGCAAAGCAGACCCTCAAACTGACACAAGACCTACAGAGAA AACCCTTTGCCAAATCTGCTCTCAGCAAGTGGACAGTGATACCGTTTACAGCTTAACACCTTTGTGAATCCCACG CCATTTCCTAACCCAGCAGAGACTGTTAATGGCCCCTTACCCTGGGTGAAGCACTTACCCTTGGAACAGAACTC TAAAAAGTATGCAAAATCTTCCTTGTACAGGGTGGTGAGCCGCCTGCCAGTGGAGGACAGCACCCCTCAGCACCA CCCACCTCATTCAGAGCACACCGTGAGCCCCGTCGGCCATTCTGTGGTGTTTTAATATTGCGATGGTTTATGG GACGTTTTAAGTGTTGTTCTTGTGTTTTCCTTTGACTTTCTGAGTTTTTCACATGCATTAACTTGCGGTAT TCCGGGTTCGAGCATCACAAGCTTTTGAGCGCATGGAACTCCATAAACTAACAAATTACATAAACTAAAGGGGGA TTTTCTTTCTTTTTTTTTTTGTTGGTAGAAAATTATCCTTTTCTAAAAACTGAACAATGGCACAATTGTTTGCTATGT GCACCCGTCCAGGACAGAACCGTGCATAGGCAAAAGGAGTGGAGCACAGCGTCCGGCCCAGTGTGTTTCCGGTTC TGAGTCAGGGTGATCTGTGGACGGGACCCCAGCACCAAGTCTACGGGTGCCAGATCAGTAGGGCCTGTGATTTCC TGTCAGTGTCCTCAGCTAATGTGAACAGTGTTGGTCTGCTGGTTAGAAACTAGAATATTGATATTTTCAGGAAAG AAAAAAAA

954/2825 FIGURE 840

MAEAPASPAPLSPLEVELDPEFEPQSRPRSCTWPLQRPELQASPAKPSGETAADSMIPEEEDDEDDEDGGGRAGS AMAIGGGGGSGTLGSGLLLEDSARVLAPGGQDPGSGPATAAGGLSGGTQALLQPQQPLPPPQPGAAGGSGQPRKC SSRRNAWGNLSYADLITRAIESSPDKRLTLSQIYEWMVRCVPYFKDKGDSNSSAGWKNSIRHNLSLHSRFMRVQN EGTGKSSWWIINPDGGKSGKAPRRAVSMDNSNKYTKSRGRAAKKKAALQTAPESADDSPSQLSKWPGSPTSRSS DELDAWTDFRSRTNSNASTVSGRLSPIMASTELDEVQDDDAPLSPMLYSSSASLSPSVSKPCTVELPRLTDMAGT MNLNDGLTENLMDDLLDNITLPPSQPSPTGGLMQRSSSFPYTTKGSGLGSPTSSFNSTVFGPSSLNSLRQSPMQT IQENKPATFSSMSHYGNQTLQDLLTSDSLSHSDVMMTQSDPLMSQASTAVSAQNSRRNVMLRNDPMMSFAAQPNQ GSLVNQNLLHHQHQTQGALGGSRALSNSVSNMGLSESSSLGSAKHQQQSPVSQSMQTLSDSLSGSSLYSTSANLP VMGHEKFPSDLDLDMFNGSLECDMESIIRSELMDADGLDFNFDSLISTQNVVGLNVGNFTGAKQASSQSWVPG

955/2825 FIGURE 841

CGGCGCGATGCGCGGAGACCCCCGCGGGGGGCGGCGGCGTGAGCCCCGATGAGGCCCCGAGCGTCCCCGGCCG ACCCCGGCCGTGCCCCCTACGTGAAGCTTGGCCTCACCGTCGTCTACACCGTGTTCTACGCGCTGCTCTTCGTG TTCATCTACGTGCAGCTCTGGCTGGTGCTGCGTTACCGCCACAAGCGGCTCAGCTACCAGAGCGTCTTCCTCTTT CTCAGCCCCTTCGTCTTCTGGCTGCTCTACTGCTTCCCTGTGTGCCTGCAGTTTTTCACCCTCACGCTGATGAAC TTGTACTTCACGCAGGTGATTTTCAAAGCCAAGTCAAAATATTCTCCAGAATTACTCAAATACCGGTTGCCCCTC TACCTGGCCTCCTCTTCATCAGCCTTGTTTTCCTGTTGGTGAATTTAACCTGTGCTGTGCTGGTAAAGACGGGA AATTGGGAGAGGAAGGTTATCGTCTCTGTGCGAGTGGCCATTAATGACACGCTCTTCGTGCTGTGTGCCGTCTCT CTCTCCATCTGTCTCTACAAAATCTCTAAGATGTCCTTAGCCAACATTTACTTGGAGTCCAAGGGCTCCTCCGTG TGTCAAGTGACTGCCATCGGTGTCACCGTGATACTGCTTTACACCTCTCGGGCCTGCTACAACCTGTTCATCCTG TCATTTTCTCAGAACAAGAGCGTCCATTCCTTTGATTATGACTGGTACAATGTATCAGACCAGGCAGATTTGAAG GTCGTTTATTTCTTCCGAGTTAGAAATCCTACAAAGGACCTTACCAACCCTGGAATGGTCCCCAGCCATGGATTC AGTCCCAGATCTTATTTCTTTGACAACCCTCGAAGATATGACAGTGATGACCTTGCCTGGAACATTGCCCCT CAGGGACTTCAGGGAGGTTTTGCTCCAGATTACTATGATTGGGGACAACAAACTAACAGCTTCCTGGCACAAGCA $\texttt{GGAACTTTGCAAGACTCAACTTTGGATCCTGACAAACCAAGCCTTGGG} \underline{\textbf{TAG}} \texttt{CATCAGTTAACAGTTTTATGGACG}$ ATTCCTCAGATGAAAAGCTTCAGAAAAGCATAGTGACAGCTGAATTTTTAGGGCACTTTTCCTTAAGAAATAGAA CTTGATTTTATTTGTTACAGGTTTCCAATGGCCCCATAGGAATAAGCAATAATGTAGACTGATAAACCCTTATT TTAGTACTAAAGAGGGAGCCTTGCTATTTCAGTGGGTATAATTTAAACTTTTTAAAGAAAATCTGTACTTTTATA ATCATGTTGTAGTTTGCACAGACTTTTATGCATAATTCACTTTAAAAATATAGAATATATGGTCTAATAGTTTTT TAAAGCTTTTGGACTAAAGTATTCCACAAATCTTACCTCTTTAGGTCACTGATGGTCACTCCGATTCTGAGTGCC ACATTGGTAGACTCCTAAAATACAGTTGACAACTTAGCCAATTGCAACTCCAGTGTTGATAATTAAAATGAAATG AAGTTTTGCCCAAAGACTGGTACTTCCTTTCAGTAGGGCGCTAATGTATACACATTAATGATAAGTTGATAACAT TAAAAATGTAGCTGACTTATCCTATTAAACCTCCTCTGCTATGTTCAC

956/2825 FIGURE 842

957/2825 FIGURE 843

GCCCGCCGGTCCACGCCGCGCACCGCTCCGAGGGCCAGCGCCACCCGCTCCGCAGCCGGCACCA<u>ATG</u>CGCGAGATC $\tt GTGCACATCCAGGCGGCCAGTGCGGCAACCAGATCGGCGCCAAGTTTTGGGAGGTCATCAGCGATGAGCATGGG$ ATCGACCCCACAGGCAGTTACCATGGAGACAGTGACTTGCAGCTGGAGAGAATCAACGTGTACTACAATGAGGCT GCTGGTAACAATATGTACCTCGGGCCATCCTGGTGGATCTGGAGCCTGGCACCATGGACTCTGTCAGGTCTGGA CCCTTCGGCCAGATCTTCAGACCAGACAACTTCGTGTTCGGCCAGAGTGGAGCCGGGAATAACTGGGCCAAGGGC CTCCAGGGCTTCCAGCTGACCCACTCTCTGGGGGGGCGCACGGGGTCCGGGATGGGCACCCTGCTCATCAGCAAG ATCCGGGAAGAGTACCCAGACCGCATCATGAACACCTTCAGCGTCATGCCCTCACCCAAGGTGTCAGACACGGTG GTGGAGCCCTACAACGCCACCCTCTCGGTCCACCAGCTGGTGGAAAACACAGATGAAACCTACTCCATTGATAAC GAGGCCCTGTATGACATCTGCTTCCGCACCCTGAAGCTGACCACCCCACCTACGGGGACCTCAACCACCTGGTG TCGGCCACCATGAGCGGGGTCACCACCTGCCTGCGCTTCCCGGGCCAGCTGAACGCAGACCTGCGCAAGCTGGCG GTGAACATGGTGCCCTTCCCTCGCCTGCACTTCTTCATGCCCGGCTTCGCGCCCCTGACCAGCCGGGGCAGCCAG CAGTACCGGGCGCTCACGGTGCCCGAGCTCACCCAGCAGATGTTCGACTCCAAGAACATGATGGCCGCCTGCGAC ATGCTCAACGTGCAGAACAGAACAGCAGCTACTTCGTGGAGTGGATCCCCAACAACGTGAAGACGGCCGTGTGC GACATCCCGCCCCGCGGCCTGAAGATGTCGGCCACCTTCATCGGCAACAGCACGGCCATCCAGGAGCTGTTCAAG $\verb|CGCATCTCCGAGCAGTTCACGGCCATGTTCCGGCGCAAGGCCTTCCTGCACTGGTACACGGGCGAGGGCATGGAC| \\$ GAGATGGAGTTCACCGAGGCCGAGAGCAACATGAACGACCTGGTGTCCGAGTACCAGCAGTACCAGGACGCCACG GCCGACGAACAAGGGGAGTTCGAGGAGGAGGAGGGCGAGGACGAGGCT**TAA**AAACTTCTCAGATCAATCGTGCAT CCTTAGTGAACTTCTGTTGTCCTCAAGCATGGTCTTTCTACTTGTAAACTATGGTGCTCAGTTTTGCCTCTGTTA GAAATTCACACTGTTGATGTAATGATGTGGAACTCCTCTAAAAAATTACAGTATTGTCTGTGAAGGTATCTATACT AATAAAAAAGCATGTGTAG

958/2825 FIGURE 844

MREIVHIQAGQCGNQIGAKFWEVISDEHGIDFTGSYHGDSDLQLERINVYYNEAAGNKYVPRAILVDLEPGTMDS VRSGPFGQIFRPDNFVFGQSGAGNNWAKGHYTEGAELVDSVLDVVRKESESCDCLQGFQLTHSLGGGTGSGMGTL LISKIREEYPDRIMNTFSVMPSPKVSDTVVEPYNATLSVHQLVENTDETYSIDNEALYDICFRTLKLTTPTYGDL NHLVSATMSGVTTCLRFPGQLNADLRKLAVNMVPFPRLHFFMPGFAPLTSRGSQQYRALTVPELTQQMFDSKNMM AACDPRHGRYLTVAAIFRGRMSMKEVDEQMLNVQNKNSSYFVEWIPNNVKTAVCDIPPRGLKMSATFIGNSTAIQ ELFKRISEQFTAMFRRKAFLHWYTGEGMDEMEFTEAESNMNDLVSEYQQYQDATADEQGEFEEEEGEDEA

959/2825 FIGURE 845

ATGGAATCAAGGGGTGTGGAATTACCTTCACTCTGGGAAAAGGCACTGAAGTTGTTGTCTGTGCTGTGAATGCCC TCGCCCACCATGTGCTCAACAAGGACCTCAAGGACATTGTTGGTGACTTCAGAGGCTTCTATAGGCAGCTCACAA GTGATGGGCAGCTCAGATGGATTGGTCCAGAAAAGGGCGTGGTGCACCTGGCGACAGCGGCCGTCCTAAACGCGG TGTGGGACTTGTGGGCCAAGCAGGAGGGAAAGCCTGTCTGGAAGTTACTTGTGGACATGGATCCCAGGATGCTGG AAATTGGTAAAAAAGAAAGAAGAAGCAAATGCTGGCACAAGGATACCCTGCTTACACGACATCGTGCGCCTGGC TGGGGTACTCAGATGACACGTTGAAGCAGCTCTGTGCCCAGGCGCTGAAGGATGGCTGGACCAGGTTTAAAGTAA AGGTGGGTGCTGATCTCCAGGATGACATGCGAAGATGCCAAATCATCCGAGACATGATTGGACCGGAAAAGACTT TGATGATGGATGCCAACCAGCGCTGGGATGTGCCTGAGGCGGTGGAGTGTCCAAGCTGGCCAAGTTCAAGC CATTGTGGATTGAGGAGCCAACCTCCCCTGATGACATTCTGGGGCACGCCACCATTTCCAAGGCACTGGTCCCAT TAGGAATTGGCATTGCCACAGGAGAACAGTGCCACAATAGAGTGATATTTAAGCAACTCCTACAGGCGAAGGCCC TGCAGTTCCTCCAGATTGACAGTTGCAGACTGGGCAGTGTCAATGAGAACCTCTCAGTATTGCTGATGGCCAAAA AGTTTGAAATTCCTGTTTGCCCCCATGCTGGTGGAGTTGGCCTCTGTGAACTGGTGCAGCACCTGATTATATTTG ACTACATATCAGTTTCTGCAAGCCTTGAAAATAGGGTGTGTGAGTATGTTGACCACCTGCATGAGCATTTCAAGT ATCCCGTGATGATCCAGCGGGCTTCCTACATGCCTCCCAAGGATCCCGGCTACTCAACAGAAATGAAGGAGGAAT GCCCCAACAACTTTTTCTTTCTGAAGTGAAAGGGCTTAAAATTTCTTGGAAATAGTTTTACAAAAATGGATTTA AAAAATCCTACCGATCAAGATGAGTTCAGCTAGAAGTCATACCACCCTCAGGAATCAGCTAAGTAATTATTACTT AGGAGGATGTTCTTACAAAAATAGCTATATCAAGGGCTGGCACCTAGACATTAAACTGTACTTTGAAAATAAAA

960/2825 FIGURE 846

MHTDPDYSAAYVVIETDAEDGIKGCGITFTLGKGTEVVVCAVNALAHHVLNKDLKDIVGDFRGFYRQLTSDGQLR WIGPEKGVVHLATAAVLNAVWDLWAKQEGKPVWKLLVDMDPRMLVSCIDFRYITDVLTEEDALEILQKGQIGKKE REKQMLAQGYPAYTTSCAWLGYSDDTLKQLCAQALKDGWTRFKVKVGADLQDDMRRCQIIRDMIGPEKTLMMDAN QRWDVPEAVEWMSKLAKFKPLWIEEPTSPDDILGHATISKALVPLGIGIATGEQCHNRVIFKQLLQAKALQFLQI DSCRLGSVNENLSVLLMAKKFEIPVCPHAGGVGLCELVQHLIIFDYISVSASLENRVCEYVDHLHEHFKYPVMIQ RASYMPPKDPGYSTEMKEESVKKHQYPDGEVWKKLLPAQEN

961/2825 FIGURE 847

 $\verb|CCTCCGGGCCTGTTGGTTGGAAGTGACGACCTTGAAGATCGGCCGGTTGGAAGTGACGACCTTGAAGATCGGCG|\\$ GGCGCAGCGGGCCGAGGGGGCGGGTCTGGCGCTAGGTCCAGCCCCTGCGTGCCGGGAACCCCAGAGGAGGTCGC TCTGCCCTACTTGATTGATGGGGCTCACAAGATCACCCAGAGCAACGCCATCCTGTGCTACATTGCCCGCAAGCA CAACCTGTGTGGGGAGACAGAAGAGGAGAAGATTCGTGTGGACATTTTGGAGAACCAGGCTATGGACGTCTCCAA TCAGCTGGCCAGAGTCTGCTACAGCCCTGACTTTGAGAAACTGAAGCCAGAATACTTGGAGGAACTTCCTACAAT GATGCAGCACTTCTCACAGTTCCTGGGGAAGAGGCCATGGTTTGTTGGAGACAAGATCACCTTTGTAGATTTCCT CGCCTATGATGTCCTTGACCTCCACCGTATATTTGAGCCCAACTGCTTGGACGCCTTCCCAAATCTGAAGGACTT CATCTCCCGCTTTGAGGGCTTGGAGAAGATCTCTGCCTACATGAAGTCCAGCCGCTTCCTCCCAAAACCTCTGTA $\texttt{CACAAGGGTGGCTGTCTGGGGCAACAAG} \underline{\textbf{TAA}} \texttt{TGCCTTGAAGGCCAGGAGGTGGGAGTGAGGAGCCCATACTCAGC}$ $\tt CTGCTGCCCAGGCTGTGCAGCGCAGCTGGACTCTGCATCCCAGCACCTGCCTCGTTCCTTTTCTCCTGTTTAT$ TCCCATCTTTACCCCCAAGACTTTATTGGGCCTCTTCACTTCCCCTAAACCCCTGTCCCATGCAGGCCCTTTGAA TGTCCCCGTGTTGCATGACAGCATTGACTGGTTTACAGGCCCTGCTCCTGCAGCATGGCCCCTGCCTTAGGCCTA

962/2825 FIGURE 848

MSMTLGYWDIRGLAHAIRLLLEYTDSSYEEKKYTMGDAPDYDRSQWLNEKFKLGLDFPNLPYLIDGAHKITQSNA ILCYIARKHNLCGETEEEKIRVDILENQAMDVSNQLARVCYSPDFEKLKPEYLEELPTMMQHFSQFLGKRPWFVG DKITFVDFLAYDVLDLHRIFEPNCLDAFPNLKDFISRFEGLEKISAYMKSSRFLPKPLYTRVAVWGNK

963/2825 FIGURE 849A

AGCCGCCCCGCTGCCGTGTCCCCTGCGGCCGGCCAGCCCCAGCCCCGGCCTCCCGCGGACCCATGCC CGCCCGTATCGGCTACTACGAGATCGACCGCACCATCGGCAAGGGCAACTTCGCGGTGGTCAAGCGGGCCACGCA CCTCGTCACCAAGGCCAAGGTTGCTATCAAGATCATAGATAAGACCCAGCTGGATGAAGAAAACTTGAAGAAGAT TTTCCGGGAAGTTCAAATTATGAAGATGCTTTGCCACCCCCATATCATCAGGCTCTACCAGGTTATGGAGACAGA ACGGATGATTTATCTGGTGACAGAATATGCTAGTGGAGGGGAAATATTTGACCACCTGGTGGCCCATGGTAGAAT GGCAGAAAAGGAGGCACGTCGGAAGTTCAAACAGATCGTCACAGCTGTCTATTTTTGTCACTGTCGGAACATTGT TCATCGTGATTTAAAAGCTGAAAATTTACTTCTGGATGCCAATCTGAATATCAAAATAGCAGATTTTGGTTTCAG AGGAAAAGAATATGATGGGCCCAAAGTGGACATCTGGAGCCTTGGAGTTGTCCTCTACGTGCTTGTGTGCGGTGC CCTGCCATTTGATGGAAGCACACTGCAGAATCTGCGGGCCCGCGTGCTGAGTGGAAAGTTCCGCATCCCATTTTT TATGTCCACAGAATGTGAGCATTTGATCCGCCATATGTTGGTGTTAGATCCCAATAAGCGCCTCTCCATGGAGCA GATCTGCAAGCACAAGTGGATGAAGCTAGGGGGACGCCGATCCCAACTTTGACAGGTTAATAGCTGAATGCCAACA ACTAAAGGAAGAAGACAGGTGGACCCCCTGAATGAGGATGTCCTCTTGGCCATGGAGGACATGGGACTGGACAA AGAACAGACACTGCAGGCGGAGCAGGCAGGTACTGCTATGAACATCAGCGTTCCCCAGGTGCAGCTGATCAACCC AGAGAACCAAATTGTGGAGCCGGATGGGACACTGAATTTGGACAGTGATGAGGGTGAAGAGCCTTCCCCTGAAGC GCAGAAGCTCCTACCTGGCTTTCCTGGAGTCAACCCCCAGGCTCCATTCCTGCAGGTGGCCCCTAATGTGAACTT GCCGCCCACGCTACAGCTGTTGAATGGAATGGGCCCCCTTGGCCGGAGGGCATCAGATGGAGGAGCCAACATCCA ACTGCATGCCCAGCAGCTGCTGAAGCGCCCACGGGGACCCTCTCCGCTTGTCACCATGACACCAGCAGTGCCAGC CTCCAACACTCTGCACCTCCCTACGGAGCGTTTCTCCCCTGTGCGCCGGTTCTCAGATGGGGCTGCGAGCATCCA GGCCTTCAAAGCTCACCTGGAAAAATGGGCAACAACAGCAGCATCAAACAGCTGCAGCAGGAGTGTGAGCAGCT GCAGAAGATGTACGGGGGGCAGATTGATGAAAGAACCCTGGAGAAGACCCAGCAGCAGCATATGTTATACCAGCA GGAGCAGCACCATCAAATTCTCCAGCAACAAATTCAAGACTCTATCTGTCCTCCTCAGCCATCTCCACCTCTTCA GGCTGCATGTGAAAATCAGCCAGCCCTCCTTACCCATCAGCTCCAGAGGTTAAGGATTCAGCCTTCAAGCCCACC CCCCAACCACCCCAACAACCATCTCTTCAGGCAGCCCAGTAATAGTCCTCCCCCCATGAGCAGTGCCATGATCCA GCCTCACGGGGCTGCATCTTCTCCCAGTTTCAAGGCTTACCTTCCCGCAGTGCAATCTTTCAGCAGCAACCTGA GAACTGTTCCTCTCCCCAACGTGGCACTAACCTGCTTGGGTATGCAGCAGCCTGCTCAGTCACAGCAGGTCAC CATCCAAGTCCAAGAGCCTGTTGACATGCTCAGCAACATGCCAGGCACAGCTGCAGGCTCCAGTGGGCGCGCAT CTCCATCAGCCCCAGTGCTGGTCAGATGCAGATGCAGCACCGTACCAACCTGATGGCCACCCTCAGCTATGGGCA $\tt CCGTCCCTTGTCCAAGCAGCTGAGTGCTGACAGTGCAGAGGGCTCACAGCTTGAACGTGAATCGGTTCTCCCCTGC$ TAACTACGACCAGGCGCATTTACACCCCCATCTGTTTTCGGACCAGTCCCGGGGTTCCCCCAGCAGCTACAGCCC $\tt TTCAACAGGAGTGGGGTTCTCTCCAACCCAAGCCCTGAAAGTCCCTCCACTTGACCAATTCCCCACCTTCCCTCC$ CAGTGCACATCAGCAGCCGCCACACTATACCACGTCGGCACTACAGCAGGCCCTGCTGTCTCCCCACGCCGCCAGA CTATACAAGACACCAGCAGGTACCCCACATCCTTCAAGGACTGCTTTCTCCCCGGCATTCGCTCACCGGCCACTC GGACATCCGGCTGCCCCCAACAGAGTTTGCACAGCTCATTAAAAGGCAGCAGCAACAACGGCAGCAGCAGCAGCA ACAGCAGCAACAGCAAGAATACCAGGAACTGTTCAGGCACATGAACCAAGGGGATGCGGGGAGTCTGGCTCCCAG CCTTGGGGGACAGAGCATGACAGAGCGCCAGGCTTTATCTTATCAAAATGCTGACTCTTATCACCACACGATCCA GAACAGCGACGATGCTTATGTACAGCTGGATAACTTGCCAGGAATGAGTCTCGTGGCTGGGAAAGCACTTAGCTC CTATCCATCTACGTGTATTACAGACATTCTGCTCAGCTACAAGCACCCCGAAGTCTCCTTCAGCATGGAGCAGGC AGGCGTGTAACAAGAAACAGAGAGAGAGCAAGAGGTCCCGAGTCCCTCCTAGTCTTTCATCCTGAATTTGCACA GAGGAAAGCGGGTGCCCGGCATGCCATCCTGATGTTGCTGGCGGGATCCCCATGCACCTTGTCCTCCACTG ATACTGGCAGCTCGGCTCCTGGACCCAAGATCCCTTGAGTGGAATTCTGCAGTGCAAGAGCCCTTCGTGGGAGCT GTCCCATGTTTCCATGGTCCCCAGTCTCCCCTCCACTTGGTGGGGTCACCAACTACTCACCAGAAGGGGGCTTAC

964/2825 FIGURE 849B

965/2825 FIGURE 850

GAAGAGTGGAGPAGRLLPPPAPGSPAAPAAVSPAAGQPRPPAPASRGPMPARIGYYEIDRTIGKGNFAVVKRATH LVTKAKVAIKIIDKTQLDEENLKKIFREVQIMKMLCHPHIIRLYQVMETERMIYLVTEYASGGEIFDHLVAHGRM AEKEARRKFKQIVTAVYFCHCRNIVHRDLKAENLLLDANLNIKIADFGFSNLFTPGQLLKTWCGSPPYAAPELFE GKEYDGPKVDIWSLGVVLYVLVCGALPFDGSTLQNLRARVLSGKFRIPFFMSTECEHLIRHMLVLDPNKRLSMEQ ICKHKWMKLGDADPNFDRLIAECQQLKEERQVDPLNEDVLLAMEDMGLDKEQTLQAEQAGTAMNISVPQVQLINP ENQIVEPDGTLNLDSDEGEEPSPEALVRYLSMRRHTVGVADPRTEVMEDLQKLLPGFPGVNPQAPFLQVAPNVNF MHNLLPMQNLQPTGQLEYKEQSLLQPPTLQLLNGMGPLGRRASDGGANIQLHAQQLLKRPRGPSPLVTMTPAVPA VTPVDEESSDGEPDQEAVQSSTYKDSNTLHLPTERFSPVRRFSDGAASIQAFKAHLEKMGNNSSIKQLQQECEQL QKMYGGQIDERTLEKTQQQHMLYQQEQHHQILQQQIQDSICPPQPSPPLQAACENQPALLTHQLQRLRIQPSSPP PNHPNNHLFRQPSNSPPPMSSAMIQPHGAASSSQFQGLPSRSAIFQQQPENCSSPPNVALTCLGMQQPAQSQQVT IQVQEPVDMLSNMPGTAAGSSGRGISISPSAGQMQMQHRTNLMATLSYGHRPLSKQLSADSAEAHSLNVNRFSPA NYDQAHLHPHLFSDQSRGSPSSYSPSTGVGFSPTQALKVPPLDQFPTFPPSAHQQPPHYTTSALQQALLSPTPPD YTRHQQVPHILQGLLSPRHSLTGHSDIRLPPTEFAQLIKRQQQQRQQQQQQQQQQQQQQEYQELFRHMNQGDAGSLAPS LGGQSMTERQALSYQNADSYHHTIQNSDDAYVQLDNLPGMSLVAGKALSSARMSDAVLSQSSLMGSQQFQDGENE ECGASLGGHEHPDLSDGSQHLNSSCYPSTCITDILLSYKHPEVSFSMEQAGV

966/2825 FIGURE 851A

 $\tt CGACCGCGGCGGCTGGAACCCTGATTGCTGTCCTTCAACGTGTTCATT{\textbf{ATG}} AAGTTATTAGTAATACTTTTGTTT$ ${\tt TCTGGACTTATAACTGGTTTTAGAAGTGACTCTTCCTCTAGTTTGCCACCTAAGTTACTACTAGTATCCTTTGAT}$ GGCTTCAGAGCTGATTATCTGAAGAACTATGAATTTCCTCATCTCCAGAATTTTATCAAAGAAGGTGTTTTGGTA GAGCATGTTAAAAATGTTTTTATCACAAAAACATTTCCAAACCACTACAGTATTGTGACAGGCTTGTATGAAGAA AGCCATGGCATTGTGGCTAATTCCATGTATGATGCAGTCACAAAGAAACACTTTTCTGACTCTAATGACAAGGAT ATGTGGCCTGGTACTGATGTACCCATTCACGATACCATCTCTTCCTATTTTATGAATTACAACTCCTCAGTGTCA TTTGAGGAAAGACTAAATAATATTACTATGTGGCTAAACAATTCGAACCCACCAGTCACCTTTGCAACACTATAT TGGGAAGAACCAGATGCAAGTGGCCACAAATACGGACCTGAAGATAAAGAAAACATGAGCAGAGTGTTGAAAAAA ATAGATGATCTTATCGGTGACTTAGTCCAAAGACTCAAGATGTTAGGGCTATGGGAAAATCTTAATGTGATCATT AAAAACTGTAGCCCTCATATGAATGTTTATCTCAAAGAAGACATTCCTAACAGATTTTATTACCAACATAATGAT CGAATTCAGCCCATTATTTTGGTTGCCGATGAAGGCTGGACAATTGTGCTAAATGAATCATCACAAAAATTAGGT GACCATGGTTATGATAATTCTTTGCCTAGTATGCATCCATTTCTAGCTGCCCACGGACCTGCATTTCACAAAGGC TACAAGCATAGCACAATTAACATTGTGGATATTTATCCAATGATGTGCCACATCCTGGGATTAAAACCACATCCC AATAATGGGACCTTTGGTCATACTAAGTGCTTGTTAGTTGACCAGTGGTGCATTAATCTCCCAGAAGCCATCGCG ATTGTTATCGGTTCACTCTTGGTGTTAACCATGCTAACATGCCTCATAATAATCATGCAGAATAGACTTTCTGTA $\texttt{CCTCGTCCATTTCTCGACTTCAGCTACAAGAAGATGATGATGATCCTTTAATTGGG\textbf{\textbf{TGA}CATGTGCTAGGGCTT}$ ATACAAAGTGTCTTTGATTAATCACAAAACTAAGAATACATCCAAAGAATAGTGTTGTAACTATGAAAAAGAATA $\verb|CTTTGAAAGAACAAGAACTTAGACTAAGCATGTTAAAATTATTACTTTGTTTTCCTTGTGTTTTTGTTTCGGTGCA|\\$ TTTGCTAATAAGATAACGCTGACCATAGTAAAATTGTTAGTAAATCATTAGGTAACATCTTGTGGTAGGAAATCA TTAGGTAACATCAATCCTAACTAGAAATACTAAAAATGGCTTTTGAGAAAAATACTTCCTCTGCTTGTATTTTGC GATGAAGATGTGATACATCTTTAAATGAAAATATACCAAAATTTAGTAGGCATGTTTTTCTAATAAATTTATATA TTTGTAAAGAAACAACAGAAATCTTTATGCAATTTGTGAATTTTGTATATTAGGGAGGAAAAGCTTCCTATATT CATTTGATAAAAGCCTGAGTTGTCACCATTATGTCTTAAGCTGTTAGTCTTAAAGATTATTGTTAAAAAATTCAG AAGAAAAGAGAGACAAGTGCTCTCTCTCTATCTATGCTTAATGCCTTTATGTAAGTTACTTAGTTGTTTGCGTG TGCCTGTGCAAGTGTGTTTGTGTGTGTGTGTGGACATTATGTGATTTACTATATAAGGAGGTCAGAGATGGAC TGTGGCCAGGCTTCCACATTCCTGAAGCACACAGATCTCAGGAAAGGTTATTTTTGCACTTCATATTTGTTTACT AACCTGAGGTATCCAAATGCTACAGAAAAATTTATGACCCAAATACAAATCTCAATTTGACTGGGACAGAATGAG GAATGGAGATTTTTGTATTTATCTTTGGGACTTTATGCCTTACTTTTTAGGCTATAGAATAGTTAAGAAATTTTTA AACAAAATTTAGTATCTTTTGGTCTTTCACACCATTCATATGTTAAGTGGCAGAATAGCCTTAGTGCTACCTCCA GTGAAGAACTGTCATCCTGCCTTTGCTAGCTGGTACCTTCTAGTAATCAAAATTAATATGAAGAAACTAGGTTGT GACAGACTAGATTATATTTAGTAGGGGAAAAATTGGGCTCAAGAACCATTCATCAGTACGTGAGACAAGCAGTTA ATAGTATGATCTTTAAAGTTTTGACAATATAAAATAAACTTGGTAACTGTTTTACAAATATAAAAGTATAATAAA TATGCAGCCCAGTTAAATATTGATTATCTGTGATGGTAAAGAACAACAGTGGTGCCAGTCATCAAACATACAGTG CGTCCTATTGAGTCACTGCTAATTTCTTGAGCCTGGTATTTGCTGCCTATTGTATTTGTGGTTGTTGAGAGGCAT CCAATAGTACTAATTTTGCTTTAAAAAAATTTCTAATTTTTTTCACATAAAACAATTATCCTAAAGGTTAATAGT TGATCGAAACAGAATAATAGAAAAATTCTACTTTAATTTCCATTAAAAAGCAAATAGCATTGACACATTTAAAGC TTTTCATTTAAAGTAGTGGATGTTTTTGAAGTATCTAAAATAGTAGCAGAATATTTTATACTTGGTCCTTGCAAT GGTGTGAGTTTTAATGATTGCATTATCGTGATTGGTGGTTATGAGTTTCAGAAATCTATACTTGGCATCCAACTC ATGAGTGGATTTTATATAGGATGGAACAGGAAGGTATGTCCTGTCAGTATCTTAACCCTTTCAACAAGACATTTA CCTATTTGTCTTTCCTTACGTTCTCAAAATATTAACTCGAATTGTAAATTAAGCAAAAATTTAAAAAGTATATGT

967/2825 FIGURE 851B

968/2825 FIGURE 852

MKLLVILLFSGLITGFRSDSSSSLPPKLLLVSFDGFRADYLKNYEFPHLQNFIKEGVLVEHVKNVFITKTFPNHY SIVTGLYEESHGIVANSMYDAVTKKHFSDSNDKDPFWWNEAVPIWVTNQLQENRSSAAAMWPGTDVPIHDTISSY FMNYNSSVSFEERLNNITMWLNNSNPPVTFATLYWEEPDASGHKYGPEDKENMSRVLKKIDDLIGDLVQRLKMLG LWENLNVIITSDHGMTQCSQDRLINLDSCIDHSYYTLIDLSPVAAILPKINRTEVYNKLKNCSPHMNVYLKEDIP NRFYYQHNDRIQPIILVADEGWTIVLNESSQKLGDHGYDNSLPSMHPFLAAHGPAFHKGYKHSTINIVDIYPMMC HILGLKPHPNNGTFGHTKCLLVDQWCINLPEAIAIVIGSLLVLTMLTCLIIIMQNRLSVPRPFSRLQLQEDDDDP LIG

969/2825 FIGURE 853

GAGATGGGTGCCCCCATCCGCACACTGTCCTTTGGCCACCGGACATCATCCCCAAGAAGAAGATGTTCCCGTGA
AGAAACCAGCAGGGCCCTCCATCTCCAAACCTGCTGCTAAGCCAGCAGCAGCAGCAGGGGGCTCCTCCAGCCAAGACCA
AAGCTGAGCCAGCTGTCCCCCAGGCCCCTCAGAAAACCCAGGAGCCTCCAGTCGATCTCTCCAAAGTGGTGATCG
AGTTTAACAAGGACCAGCTGGAGGAGTTCAAGGAGGCCTTCGAGCTGTTTGACCGAGTGGGGGATGGCAAGATCC
TGTACAGCCAGTGTGGGGACGTGATGAGGGGCCCTGGGCCAGAACCCCACCAACGCCGAGGTGCTCAAGGTCCTGG
GGAACCCCAAGAGTGATGAGCTGAAGTCGCGGCGTGTGGACTTTTGAGACTTTCCTGCCCATGCTCCAGGCAGTGG
CCAAGAACCGAGGCCAAGGCACATATGAGGACTACTTGGAGGGGTTTCGTGTGTTTTGACAAGGAGGGGAACCGC
AAGTCATGGGAGCAGAGCTCAGACATGTTCTCACCACCCTTGGAGAGAAGATGACTGAGGAGGAGGGGAGACCG
TTCTGGCAGGACACGAGGACAGCAACGGCTGCATCAACTACGAGGCCTTCTTGAAACACATCCTAAGCGTCTGAG
TGCTGCAGATCCAGTGGGGTCCGGACACTGGGCCCCGCAGGCGAAAGCACGTTCCAGCCACCAGGAGGCCACCTA
TTGTTTCAAAATAAAGACTGGGTTCCTC

970/2825 FIGURE 854

 ${\tt MPPKKDVPVKKPAGPSISKPAAKPAAAGAPPAKTKAEPAVPQAPQKTQEPPVDLSKVVIEFNKDQLEEFKEAFEL} \\ {\tt FDRVGDGKILYSQCGDVMRALGQNPTNAEVLKVLGNPKSDELKSRRVDFETFLPMLQAVAKNRGQGTYEDYLEGF} \\ {\tt RVFDKEGNGKVMGAELRHVLTTLGEKMTEEEVETVLAGHEDSNGCINYEAFLKHILSV} \\ {\tt MPPKKDVPVKKPAGPSISKPAAKPAAAGAPPAKTKAEPAVPQAPQKTQEPPVDLSKVVIEFNKDQLEEFKEAFEL} \\ {\tt FDRVGDGKVMGAELRHVLTTLGEKMTEEEVETVLAGHEDSNGCINYEAFLKHILSV} \\ {\tt MPPKKDVPVKKPAGPSISKPAAKPAAAGAPPAKTKAEPAVPQAPQKTQEPPVDLSKVVIIEFNKDQLEEFKEAFEL} \\ {\tt FDRVGDGKVMGAELRHVLTTLGEKMTEEEVETVLAGHEDSNGCINYEAFLKHILSV} \\ {\tt MPPKKDVPVKMGAELRHVLTTLGEKMTEEEVETVLAGHEDSNGCINYEAFLKHILSV} \\ {\tt MPPKKDVPVKMGAELRHVLTTLGEKMTEEEVETVLAGHEDSNGCINYEAFLKHILSV} \\ {\tt MPPKKDVPVKMGAELRHVLTTLGEKMTEEEVETVLAGHEDSNGCINYEAFLKHILSV} \\ {\tt MPPKKDVPVKMGAELRHVLTTLGEKMTEEEVETVLAGHEDSNGCINYEAFLKHILSV} \\ {\tt MPPKMGAELRHVLTTLGEKMTEEEVETVLAGHEDSNGCINYEAFLKHILSV} \\ {\tt MPPKMGAELRHVLTTLGEKMTEETVLAGHEDSNGCINYEAFLKHILSV} \\ {\tt MPPKMGAELRHVLTTLGEKMTEETVLAGHETVLAGHEDSNGCINYEAFLKHILSV} \\ {\tt MPPKMGAELRHVLTTLGEKMTEETVLAGH$

971/2825 FIGURE 855

972/2825 FIGURE 856

 $\label{thm:modetvgnvvllaivtlisvvqngffahkvehesrtqngrsfqrtgtlafervytanqncvdayptflavlwsagllcsqvpaafaglmylfvrqkyfvgylgertqstpgylfgkriilflflmsvagifnyylifffgsdfenyiktisttisplllip$

973/2825 FIGURE 857

974/2825 FIGURE 858

MSAQESCLSLIKYFLFVFNLFFFVLGSLIFCFGIWILIDKTSFVSFVGLAFVPLQIWSKVLAISGIFTMGIALLG CVGALKELRCLLGLYFGMLLLLFATQITLGILISTQRAQLERSLRDVVEKTIQKYGTNPEETAAEESWDYVQFQL RCCGWHYPQDWFQVLILRGNGSEAHRVPCSCYNLSATNDSTILDKVILPQLSRLGHLARSRHSADICAVPAESHI YREGCAQGLQKWLHNNLISIVGICLGVGLLELGFMTLSIFLCRNLDHVYNRLARYR

975/2825 FIGURE 859

CCGAGGCGACAGCTGTGGCCGGAAGCGGGCTGTTGGCGGCTGCCTGGCCAAAGACGGCTTGCAGCAGTCTAAGT GCCCGGACACTACCCCAAAACGGCGCGCGCCTCGTCGCTGTCGCGTGACGCCGAGCGCCGAGCCTACCAATGGT GCCGGGAGTACTTGGGCGGGGCCTGGCGCCGAGTGCAGCCCGAGGAGCTGAGGGTTTACCCCGTGAGCGGAGGCC TCAGCAACCTGCTCTCCGCTGCTCCCGGACCACCTGCCCAGCGTTGGCGAGGAGCCCCGGGAGGTGCTTC TGCGGCTGTACGGAGCCATCTTGCAGGGCGTGGACTCCCTGGTGCTAGAAAGCGTGATGTTCGCCATACTTGCGG AGCGGTCGCTGGGGCCCCAGCTGTACGGAGTCTTCCCAGAGGGCCGGCTGGAACAGTACATCCCAAGTCGGCCAT TGAAAACTCAAGAGCTTCGAGAGCCAGTGTTGTCAGCAGCCATTGCCACGAAGATGGCGCAATTTCATGGCATGG AGATGCCTTTCACCAAGGAGCCCCACTGGCTGTTTGGGACCATGGAGCGGTACCTAAAACAGATCCAGGACCTGC $\verb|CCCCAACTGGCCTCCCTGAGATGAACCTGCTGGAGATGTACAGCCTGAAGGATGAGATGGGCAACCTCAGGAAGT|\\$ TACTAGAGTCTACCCCATCGCCAGTCGTCTTCTGCCACAATGACATCCAGGAAGGGAACATCTTGCTGCTCTCAG AGCCAGAAAATGCTGACAGCCTCATGCTGGTGGACTTCGAGTACAGCAGTTATAACTATAGGGGCTTTGACATTG GGAACCATTTTTGTGAGTGGGTTTATGATTATACTCACGAGGAATGGCCTTTCTACAAAGCAAGGCCCACAGACT ACCCCACTCAAGAACAGCAGTTGCATTTATTCGTCATTACCTGGCAGAGGCAAAGAAGGTGAGACCCTCTCCC GGGGTCTGTGGTCCATCCTCCAGGCATCCATGTCCACCATAGAATTTGGTTACTTGGACTATGCCCAGTCTCGGT TCCAGTTCTACTTCCAGCAGAAGGGGCAGCTGACCAGTGTCCACTCCTCATCCTGACTCCACCCTCCCACTCCTT AGAGCCCCGTGCTGTGTATGTAACACAATAAACAAGCTTC

976/2825 FIGURE 860

MAAEATAVAGSGAVGGCLAKDGLQQSKCPDTTPKRRRASSLSRDAERRAYQWCREYLGGAWRRVQPEELRVYPVS GGLSNLLFRCSLPDHLPSVGEEPREVLLRLYGAILQGVDSLVLESVMFAILAERSLGPQLYGVFPEGRLEQYIPS RPLKTQELREPVLSAAIATKMAQFHGMEMPFTKEPHWLFGTMERYLKQIQDLPPTGLPEMNLLEMYSLKDEMGNL RKLLESTPSPVVFCHNDIQEGNILLLSEPENADSLMLVDFEYSSYNYRGFDIGNHFCEWVYDYTHEEWPFYKARP TDYPTQEQQLHFIRHYLAEAKKGETLSQEEQRKLEEDLLVEVSRYALASHFFWGLWSILQASMSTIEFGYLDYAQ SRFQFYFQQKGQLTSVHSSS

977/2825 FIGURE 861

 $\tt CCGGCTCGGAGTGTGATCTAAGCAGGTCGCGTACCTTCCTCAGGTGACTCCGGCCACAGCCCATTGTCCGCGGCC$ ACCGGCGGAGTTTAGCCGCAGACCTCGAAGCGCCCCGGGGTCCTTCCCGAACGGCAGCGGCTGCGGCGGGTCCAT **G**GAGAAGGGCCCTGTGCGGGCACCGGCGGAGAAGCCGCGGGGCGCCAGGTGCAGCAATGGGTTCCCCGAGCGGGA TCCGCCGCGGCCCGGGCCCAGCAGGCCGGCGGAGAAGCCCCCGCGGCCCAAGAGCGCGCAGCCCGCGGA CGGCTGGAAGGGCGAGCGCCCCGCAGCGAGGAGGATAACGAGCTGAACCTCCCTAACCTGGCAGCCGCCTACTC GTCCATCCTGAGCTCGCTGGGCGAGAACCCCCAGCGGCAAGGGCTGCTCAAGACGCCCTGGAGGGCGGCCTCGGC CATGCAGTTCTTCACCAAGGGCTACCAGGAGACCATCTCAGATGTCCTAAACGATGCTATATTTGATGAAGATCA TGATGAGATGGTGATTGTGAAGGACATAGACATGTTTTCCATGTGTGAGCATCACTTGGTTCCATTTGTTGGAAA GGTCCATATTGGTTATCTTCCTAACAAGCAAGTCCTTGGCCTCAGCAAACTTGCGAGGATTGTAGAAATCTATAG TAGAAGACTACAAGTTCAGGAGCGCCTTACAAAACAAATTGCTGTAGCAATCACGGAAGCCTTGCGGCCTGCTGG AGTCGGGGTAGTGGTTGAAGCAACACACATGTGTATGGTAATGCGAGGTGTACAGAAAATGAACAGCAAAACTGT AGTCAAATAAATATAATAAAGGGTTGAGCCTCTACTTTCTTCTTGCCACCTTTTTGTGGCAATATTAAAGTGAAC TGCTAATAGTGTAAGTATGTGCACAAAACCACTGCCAGATAACCAGAGGGGCCTGGGAAGGAGAAGAATTAGTG TATTTTTTCAAATAGTACAGTAATTTGCCTCATAAGCATAGGAGCATTGGGAATGAGAGGGAACTGTGCCCAGT ATACTGTTTTTTTCTTCCTCCAATAAAAGTGGTGTAGTGCCGAAAGTGCTAAAATATTTAGTGCGGTATTGCTC TGTGAATTCAAGTTCAACAGACTTCACTTTGGTCATGTTTATTAAACCACCAGTGACATTTAAAAATATATTTTT AGCAGTCGTAATGTTAGTCACCAAGGGAAGGTGGTGGAATGTCTATGTTTTTGATTTTACTGTGAGTTAAAAAGG CACATTTCTACCTTCTATTGTTTTTAAATTCAAGAATAGGGAATTAGTTCCTGGTGTTGTTTACGAGTGTATTCT CGTGTCAACATACAGGGATTTAGACATTTAACTCTCTGTGCCTTGATAAGAATATCATTTAGAGTGTAGATACTT TTGCCTTTTTAAAAAAGCCATTATTTTATGAGACTTAGTACTCACACTGCAAATAACTAGTCAGCTCAGTTTTAA CTTTATAGGTTTATTGAGTTTCCTTTGTGTGATCCATGTAGATGCCTCAAAATGTTTCTTCTTCTTCTTTTTTT AATCTTATAAGATATTTTCTAAGTATTTCCAGAAACATTTGAGAGTGCCCATCATTTTCAGGTCTGCAGAACCA TAGCTTCCACGCACCTGAACGAGCACAGAATGAACTGACGGTGGAAGACATTATGAGCTGTGTCCAACGTTTTAA CCAAAGCGTATCGTACCAACGATCTGTGAAAATGCACTGGAAGCTTCTGGTCCCGGTTTCCTTTGTGGTCTATGT GGGTCTTGTCCTCATTGTAACTCCGTATAGATGGTATAGGTATTTTAATCCTGGAAGCTGTTGCCTTATTAATGA CACAAACACAGTTATTCCATGAAGTTTAGTATTTGGTTGACATAGTGCTCTTCAAATTCATCCCATTACCCTAAA AGTAATAACTTTGATGCTTTAACTTTAGTCCCATCTCTGCCACTTTGATGCTATTTGGGTTATGATGGGGC AAGATGCCAGAGGTATTGGGTTTTTTTTTTTTTCCATTCCTCTACTTCTGTTTCCTAGCTTTTTCTTCTG GAGTTTAAGTACAGTGATGGTTGGCTTGAGTACCTTTTTAAATCTAGCCCAGTATAAACATTAGCCTGCTTAATA GAGGACCAAAGAAATTGTCAGCTATACATTTATCTTTATGAACTCATTTATATTCCTTTTTAATGACTCGTTGTT CTAACATTTCCTAGAAGTGTTCTTATAAAGGTCTAATGTATCCACAGGCTGTTGTCTTATTAGTAAATGCAAAGT AATGACTTTGTCTGTTTTACTCTAGTCTTTAGTACTTCAAAATTACCTTTTCATATCCATGATCTTGAGTCCATT

978/2825 FIGURE 862

MEKGPVRAPAEKPRGARCSNGFPERDPPRPGPSRPAEKPPRPEAKSAQPADGWKGERPRSEEDNELNLPNLAAAY SSILSSLGENPQRQGLLKTPWRAASAMQFFTKGYQETISDVLNDAIFDEDHDEMVIVKDIDMFSMCEHHLVPFVG KVHIGYLPNKQVLGLSKLARIVEIYSRRLQVQERLTKQIAVAITEALRPAGVGVVVEATHMCMVMRGVQKMNSKT VTSTMLGVFREDPKTREEFLTLIRS

979/2825 **FIGURE 863**

980/2825 FIGURE 864

 ${\tt MAAAMVPGRSESWERGEPGRPALYFCGSIRGGREDRTLYERIVSRLRRFGTVLTEHVAAAELGARGEEAAGGDRLIHEQDLEWLQQADVVVAEVTQPSLGVGYELGRAVAFNKRILCLFRPQSGRVLSAMIRGAADGSRFQVWDYEEGEVEALLDRYFEADPPGQVAASPDPTT}$

981/2825 FIGURE 865

982/2825 FIGURE 866

MQKYEKLEKIGEGTYGTVFKAKNRETHEIVALKRVRLDDDDEGVPSSALREICLLKELKHKNIVRLHDVLHSDKK LTLVFEFCDQDLKKYFDSCNGDLDPEIVKSFLFQLLKGLGFCHSRNVLHRDLKPQNLLINRNGELKLADFGLARA FGIPVRCYSAEVVTLWYRPPDVLFGAKLYSTSIDMWSAGCIFAELANAGRPLFPGNDVDDQLKRIFRLLGTPTEE QWPSMTKLPDYKPYPMYPATTSLVNVVPKLNATGRDLLQNLLKCNPVQRISAEEALQHPYFSDFCPP

983/2825 FIGURE 867A

TCTTTAGAGCAAAAGGTGAATCAAAGGAGAGTCCCAGAAAACCTGTGACTGTTGAAGAAAATTCATCTGTGAATT TTTATATTCAAGGAGTCAGTATTTATATTCATCTTTTAAACTGGGAAGATTTATATTTTACTTTAAAACTTCTTG ATAATAATTTACA**ATG**AATGGACACAGTGATGAAGAAAGTGTTAGAAACAGTAGTGGAGAATCAAGCCAGTCGGA TGATGATTCTGGGTCAGCTTCAGGCTCTGGATCTGGTTCGAGTTCTGGAAGCAGTAGTGATGGAAGCAGTAGCCA GTCAGGTAGCAGTGACTCTGACTCCGGATCTGAATCAGGCAGTCAGAGTCTGAGTCAGACACTTCCCGAGA AAACAAAGTTCAAGCAAAACCACCGAAAGTTGATGGAGCTGAGTTTTGGAAATCTAGTCCTAGTATTCTGGCCGT TCAGAGATCTGCAATCCTCAAGAAGCAGCAACAGCAGCAGCAGCAACAACATCAAGCCTCATCTAATAGCGG ATCAGAAGAGGATTCCTCTAGCAGTGAAGATTCCGATGACTCATCAAGTGAGGTCAAAAGGAAAAAGCATAAAGA GAAAAGCAGTTGTGATGAAACAGAATCTGATTATGAGCCAAAAAACAAAGTCAAAAGCAGAAAACCTCAAAATAG TGAAGAAGATTATGATAATGATAAAAGAAGTTCTCGTCGCCAAGCAACTGTTAATGTTAGCTATAAGGAGGATGA AGAAATGAAAACAGATTCTGATGACCTACTGGAAGTCTGTGGAGAGGATGTTCCTCAACCTGAGGAAGAGGAATT ATGGAAAGGATGGTCCCATATCCACAACACTTGGGAGACAGAAGAAACCCTCAAGCAGCAGAATGTTAGAGGAAT GAAAAAATTGGATAATTATAAGAAAAAAGATCAGGAAACAAAAAGATGGTTGAAAAATGCCTCTCCAGAAGATGT GGAATATTATAATTGCCAGCAAGAACTTACAGATGATCTACATAAACAGTATCAAATAGTGGGACGTATAATTGC AAAAACCACTCCTTTTAAAGATTGCAAAGTATTAAAACAAAGGCCAAGGTTTGTAGCCCTGAAGAAGCAGCCATC CTATATTGGAGGACATGAGGGCTTAGAATTAAGAGATTATCAACTGAATGGTTTAAATTGGCTTGCTCATTCTTG GTGCAAAGGAAATAGTTGCATACTCGCTGATGAAATGGGCCTTGGAAAAACAATACAGACGATCTCATTTCTGAA TTATTTGTTTCATGAACATCAATTATATGGACCTTTTTTATTGGTAGTACCGCTCTCCACTCTTACTTCCTGGCA AAGAACTCATGAATGGACGCATCATCAGACCAAACGGTTAAAATTTAATATATTGTTAACAACTTATGAAATTTT ATTAAAAGATAAGGCATTCCTTGGAGGTCTAAATTGGGCATTTATAGGTGTTGATGAAGCACCCGATTAAAGAA TGATGACTCCCTTCTGTATAAAACTTTAATAGATTTTAAATCCAATCATCGTCTCCTTATCACTGGAACTCCTCT ACAGAATTCCCTCAAAGAGCTCTGGTCTTTGCTACATTTCATTATGCCAGAAAAGTTTTCTTCCTGGGAAGATTT CCGAGTTAAGAAAGATGTGGAAAAATCTCTTCCTGCCAAGGTTGAGCAGATTTTAAGAATGGAAATGAGTGCTTT ACAGAAACAATATTACAAATGGATTTTAACTAGGAATTACAAAGCCCTCAGCAAAGGTTCCAAGGGCAGTACCTC AGGCTTTTTGAACATTATGATGGAGCTAAAGAAATGTTGTAACCATTGCTACCTCATTAAACCACCAGATAATAA TGAATTCTATAATAAACAGGAGGCCTTACAACACTTAATTCGTAGTAGCGGAAAATTGATTCTTCTTGACAAGCT ATTAATTCGCCTAAGAGAACGAGGCAATCGAGTTCTTATTTTTTCACAAATGGTGCGGATGTTAGATATACTTGC AGAATATTTGAAATATCGTCAATTCCCCTTTCAAAGATTAGATGGATCAATAAAAGGAGAACTGAGGAAACAAGC TCTAGATCATTTTAATGCTGAGGGATCAGAGGATTTTTGCTTTTTGCTGTCCACAAGAGCTGGAGGTCTAGGGAT TAATTTAGCCTCTGCTGACACTGTTGTTATATTTGATTCCGATTGGAATCCACAGAATGATCTTCAGGCACAGGC TAGAGCCCATCGAATTGGCCAAAAGAAACAGGTGAATATTTATCGTCTAGTTACAAAGGGATCAGTTGAAGAAGA TATTCTTGAAAGGGCGAAAAAGAAGATGGTTTTAGATCATCTTGTAATTCAAAGAATGGACACAACTGGGAAGAC AGTACTACATACAGGTTCTGCCCCATCAAGTTCTACTCCTTTCAATAAAGAAGAGTTATCAGCCATTTTAAAGTT TGGTGCTGAAGAACTTTTTAAGGAACCTGAAGGAGAAGAACAAGAGCCCCAGGAAATGGATATAGATGAAATCTT GAAGAGAGCTGAAACTCATGAAAATGAACCAGGTCCTTTAACTGTAGGAGATGAATTGCTTTCCCAGTTCAAGGT TGCCAACTTCTCAAATATGGATGAGGATGACATTGAGTTGGAACCTGAAAGAATTCAAAGAATTGGGAGGAAAT TATTCCAGAAGATCAAAGAAGACGATTAGAAGAAGAAGAAAAGACAAAAGGAACTTGAAGAAATTTATATGCTCCC AAGAATGAGAAATTGTGCAAAACAGATTAGTTTCAATGGAAGTGAAGGGAGGCGCAGTAGAAGTAGGAGATACTC TGGATCTGATAGTGATTCCATCTCAGAAGGGAAAAGGCCAAAGAAACGTGGAAGACCACGGACTATTCCTCGGGA GAATATTAAAGGATTTAGTGATGCAGAAATTAGGCGGTTTATCAAGAGCTATAAGAAATTTGGTGGTCCTCTGGA

984/2825 **FIGURE 867B**

AAGATTAGATGCAATTGCTCGAGATGCTGAGTTAGTTGATAAGTCAGAAACAGACCTTAGACGACTGGGAGAATT AGTGAAGGGTCCAACATTCCGAATATCAGGAGTACAGGTGAATGCCAAACTAGTCATCTCCCATGAAGAAGAATT AATACCTTTGCACAAATCCATTCCTTCTGATCCAGAAGAAGAAGCAGTATACTATCCCATGCCACACAAAGGC AGCTCATTTTGATATAGACTGGGGCAAAGAAGATGATTCCAATTTGTTAATTGGCATCTATGAATATGGATATGG AAGCTGGGAAATGATTAAAATGGATCCTGACCTCAGTCTAACACACAAGATTCTTCCAGATGATCCCGATAAAAA ACCACAAGCAAAACAGTTGCAGACCCGTGCAGACTACCTCATCAAATTACTTAGTAGAGATCTTGCAAAAAAAGA AGCTCTTTCTGGTGCGGGAAGTTCAAAGAGGAGAAAAGCAAGAGCTAAGAAGAATAAAGCAATGAAGTCTATAAA AGTGAAAGAGGAAATAAAGAGTGATTCTTCTCCTCTGCCTTCAGAGAAGTCTGATGAAGATGATGATAAATTGAG TGAATCCAAGTCTGATGGTAGGGAAAGATCCAAGAAATCTTCAGTGTCAGATGCTCCAGTTCATATCACGGCAAG GAGGCCTGTTAAAGCAGCTTTGAAACAACTTGATAGGCCTGAGAAAGGCCTTTCAGAAAGAGAACAACTAGAACA TACTAGACAATGTTTAATAAAAATTGGAGACCATATCACAGAATGTCTAAAAGAGTATACAAATCCTGAACAAAT TAAGCAATGGAGAAAAACCTGTGGATTTTTGTATCTAAGTTTACTGAATTTGATGCAAGAAAATTACATAAATT ATATAAGCATGCTATTAAAAAACGGCAGGAGTCTCAGCAAAACAGTGATCAAAACAGCAACTTGAATCCTCACGT GATTAGAAATCCAGATGTGGAAAGATTAAAAGAGAATACAAATCACGATGATAGCAGCAGGGACAGTTATTCCTC TGATAGACACTTAACTCAGGATCCATGATCATAAAGACCGACATCAGGGAGATTCTTACAAAAAAGTGATTC ATATTACAGTGACAGAGAAACACAGAAAACTGGATGATCACAGGAGTAGAGATCACAGGTCAAATTTGGAAGG TTCTGAATATACGCACCATAAATCTTCCAGGGATTATAGGTATCACTCAGACTGGCAAATGGACCACAGAGCTTC CAGCAGTGGCCCTAGGTCACCACTAGATCAGAGATCTTATGGCTCCAGATCTCCATTTGAACATTCAGTTGAACA ${ t CAAAAGTACACCGGAGCATACCTGGAGTAGTCGGAAAACA}{ t TAA}{ t CAAAAACTGATACTTCGTCTTTCTGGACTTTT}$ CTTTTAGCCATATATCATAAACCAACACAGTAATTGCCTTACATGACTGAAAGATATAAACAGATCTTCTATCA GTAGCAGTATTGTTACTTCTTTCCAGGATGCAAGGTCTATTATCCCAACAGAAGAGAAAATATTTTTATATTTAA CTATTACAGGGACCTCAACACTGCCCTCCCATACAGGCTGGATAAAACTGTTTTTAAGTCAGTGATTTTAGACTG GTTCTTTCTTAATAAAGAAAAACACTCAAGGACTTTGTTCATTTCCAAAGCTACTTGTTTACATTGTACACTGCG ACCACCTTGCCGCTTTTCATCACAAGCTTGAATATTTAAATTCTGTACTTATATCTGTAAAATAGCCAGGAATTT CCTGTTTGTGATCTATTATGCCTTTTTACAAAAAAATGGCTGTAAATTATTGTAAATATTAAAGGAACTTTCCT TACAAAAAAAAAAAAAA

985/2825 FIGURE 868

MNGHSDEESVRNSSGESSQSDDDSGSASGSGSGSSSSGSSSDGSSSSDSDSGSESGSQSESESDTSRENKVQ AKPPKVDGAEFWKSSPSILAVQRSAILKKQQQQQQQQQHQASSNSGSEEDSSSSEDSDDSSSEVKRKKHKDEDWQ MSGSGSPSQSGSDSESEEEREKSSCDETESDYEPKNKVKSRKPQNRSKSKNGKKILGQKKRQIDSSEEDDDEEDY DNDKRSSRRQATVNVSYKEDEEMKTDSDDLLEVCGEDVPQPEEEEFETIERFMDCRIGRKGATGATTTIYAVEAD GDPNAGFEKNKEPGEIQYLIKWKGWSHIHNTWETEETLKQQNVRGMKKLDNYKKKDQETKRWLKNASPEDVEYYN CQQELTDDLHKQYQIVGRIIAHSNQKSAAGYPDYYCKWQGLPYSECSWEDGALISKKFQACIDEYFSRNQSKTTP FKDCKVLKQRPRFVALKKQPSYIGGHEGLELRDYQLNGLNWLAHSWCKGNSCILADEMGLGKTIQTISFLNYLFH EHQLYGPFLLVVPLSTLTSWQREIQTWASQMNAVVYLGDINSRNMIRTHEWTHHOTKRLKFNILLTTYEILLKDK AFLGGLNWAFIGVDEAHRLKNDDSLLYKTLIDFKSNHRLLITGTPLQNSLKELWSLLHFIMPEKFSSWEDFEEEH GKGREYGYASLHKELEPFLLRRVKKDVEKSLPAKVEQILRMEMSALQKQYYKWILTRNYKALSKGSKGSTSGFLN IMMELKKCCNHCYLIKPPDNNEFYNKQEALQHLIRSSGKLILLDKLLIRLRERGNRVLIFSQMVRMLDILAEYLK YRQFPFQRLDGSIKGELRKQALDHFNAEGSEDFCFLLSTRAGGLGINLASADTVVIFDSDWNPONDLQAQARAHR IGQKKQVNIYRLVTKGSVEEDILERAKKKMVLDHLVIQRMDTTGKTVLHTGSAPSSSTPFNKEELSAILKFGAEE LFKEPEGEEQEPQEMDIDEILKRAETHENEPGPLTVGDELLSQFKVANFSNMDEDDIELEPERNSKNWEEI1PED QRRRLEEEERQKELEEIYMLPRMRNCAKQISFNGSEGRRSRSRRYSGSDSDSISEGKRPKKRGRPRTIPRENIKG FSDAEIRRFIKSYKKFGGPLERLDAIARDAELVDKSETDLRRLGELVHNGCIKALKDSSSGTERTGGRLGKVKGP TFRISGVQVNAKLVISHEEELIPLHKSIPSDPEERKQYTIPCHTKAAHFDIDWGKEDDSNLLIGIYEYGYGSWEM IKMDPDLSLTHKILPDDPDKKPQAKQLQTRADYLIKLLSRDLAKKEALSGAGSSKRRKARAKKNKAMKSIKVKEE IKSDSSPLPSEKSDEDDDKLSESKSDGRERSKKSSVSDAPVHITASGEPVPISEESEELDQKTFSICKERMRPVK AALKOLDRPEKGLSEREOLEHTROCLIKIGDHITECLKEYTNPEOIKOWRKNLWIFVSKFTEFDARKLHKLYKHA IKKRQESQQNSDQNSNLNPHVIRNPDVERLKENTNHDDSSRDSYSSDRHLTQYHDHHKDRHQGDSYKKSDSRKRP YSSFSNGKDHRDWDHYKQDSRYYSDREKHRKLDDHRSRDHRSNLEGSLKDRSHSDHRSHSDHRLHSDHRSSSEYT HHKSSRDYRYHSDWQMDHRASSSGPRSPLDQRSYGSRSPFEHSVEHKSTPEHTWSSRKT

986/2825 FIGURE 869

CCGGAGCGGGGATCACCTTAACTTGACTGTGGTCCCCCAGCCCTCCAAGAAATGCCCATAAACCAAGGAGGAGAG GGTAAAAAATAGAAGAATACATGGAATTGGGGAGAAGAATGGGAAAAAAAGATGGGAGTGACTGCATAATGTCAGC ATTTTGTGCTTTTGGCTCAGCATTTGGATTGGATGGAGGATTTTCATCCTGAAACCAGCCCTCCGTCGCCACCGA CCATGGAATAATTGTCTTCCACGAAACTCTTCCCTGGTGCCAAAAAGGGTTTTATGTATATTTTGAGAGCTCCA ${\tt CATCCCTGGATCTGAATCCTCCACTTCCCACTGGAACCATGCCCCTCCCAGTCCCGTTTCTCCCCTCTAGGGGGAT}$ $\underline{\mathbf{G}}$ GAGGCTGAGAGACCCCAGGAAGAAGAGGATGGTGAGCAGGGCCCCCTCAGGATGAGGAAGGCTGGCCCCCTCC AAACTCCACCACTCGGCCTTGGCGATCTGCTCCTCCATCCCTCCTCCAGGGACCCGCCACACAGCCCTGGG ACCCCGCTCGGCCTCCCTGCTCCCTGCAGACTGAACTCCTTCTGGACCTGGTGGCTGAAGCCCAGTCCCGCCG CCTGGAGGAGCAGAGGCCACCTTCTACACCCCCCAAAACCCCTCAAGCCTAGCCCCTGCCCCACTCCGTCCTCT CGAGGACAGAGAACAGCTTTACAGCACTATCCTCAGTCACCAGTGCCAGCGGATGGAAGCCCAGCGGTCAGAGCC TCCCCTCCCAGGGGGGCAAGAGCTCCTGGAGTTGCTGCTGAGAGTTCAGGGTGGGGGTCGAATGGAGGAGCA $\mathtt{AAGGTCCCGGCCCCCACACACCTGC}$ TGGGCAGCCCATTGCATGCCCTCAACTCTTGCTTGGCAGGGGTACCAGAGACTGAAAGACACGGCACAAATCTCA ATATTCATCTCCCACATCACCTTCCCTGGGAACTGGACAGGGTGAAAGTCCTCAAACTCTGGGAACAGGCGAGAT GGAACAGGGATTTAACTCCCCGCCCACAGGTCCATGGGAGCTTGAGGCAGTAAGGGGGATCCCAGGCACCCATCT CAAGGAGTGGCTGGGAGTCTTTTCCCTAACTTGTGGGGACACCACCAGTTGTCAAGCTACTAGGCAGTAGGGTCT GAGGGCTCAGGCCTCCACCTGAGAGGGTTATAACCTGAGAGACAGCTCTACCCTTCCTCCCAGTAAGAAGGGAAGG TGGGTGGGCACCTGAGAGATTAAGACTATTCTCCCAGTCCCACTACCAGCACCCCCGATCCCTGAGACTGAGGGG

987/2825 FIGURE 870

 ${\tt MEAERPQEEEDGEQGPPQDEEGWPPPNSTTRPWRSAPPSPPPPGTRHTALGPRSASLLSLQTELLLDLVAEAQSR} \\ {\tt RLEEQRATFYTPQNPSSLAPAPLRPLEDREQLYSTILSHQCQRMEAQRSEPPLPPGGQELLELLLRVQGGGRMEE} \\ {\tt QRSRPPTHTC} \\$

988/2825 FIGURE 871

TTCGTGTTCAGCTCGCGTCCTGCAGCTGTCCGAGGTGCTCCAGTTGGAGGCTGAGGTTCCCGGGCTCTGTCGCTG CAGCAGAGCTGGAAGTCGAGTGTGCTACTCAACTCAGGAGATTTGGAGACAAACTGAACTTCCGGCAGAAACTTC ${\tt TGAATCTGATATCCAAACTCTTCTGCTCAGGAACC} {\tt TGA} {\tt CTGCATCAAAAACTTGCATGAGGGGACTCCTTCAAAA}$ GAGTTTTCTCAGGAGGTGCACGTTTCATCAATTTGAAGAAGACTGCATTGTAATTGAGAGGAATGTGAAGGTGC ATTCATGGGTGCCCTTGGAAACGGAAGATGGAATACATCAAAGTGAATTTCTGTTCAAGTTTTCCCAGATTATCA TTCTTTGGGATGAGAGACATTATAAAACCACTTTGTTTATTTTAAAGCAAGAATGGAAGACCCTTGAAAATAAA GAAGTAATTATTGACACATTTCTTTTTTACTTAGAGAATCGTTCTAGTGTTTTTTGCCGAAGATTACCGCTGGCCT ACTGTGAAGGTAGATGACCTGTGATTAGACTGGGCGGCTGGGGAGAAACAGTTCAGTGCATTGTTGTTGCTG TTTTTGGTGTTTTCAGTGCCAACTCAGCACATTGTATATGATTCGGTTTATACATATTACCTTGTTATA ATGAAAAAACTCATTCTGAGAACACTGAAATGTTATACTCAGTGTTGATTTCTTCGGTCACTACACAACGTAAAA TTACTAGAAACTATGGAAAACTGGAAAATAACTTTGAAAAAATTGGATAAAGTATAGGAGGGTTACTTGGGGCCA GTAAATCAGTAGACTGAACATTCAATATAATAAAAGAACATGGGGATTTTGTATAACCAGGGATAATAAAAAGAA AAAGAAGTTAATTTTAATTGATGTTTTTGAAACTTAGTAGAACAAATATTCAGAAGTAACTTGATAAGATATGA ATGTTTCTAAAGAGTTTCTAAAGGTTCGAAATGCTCCTTGTCACATTAGTGTGCATCCTACAAAAAGTGATCTCT AATGAAAGTTCCATATATGCATTGGTGAATATATGTATACACATACTTACATACTTATATGGGTATCTGTATA AAAGTATTTTCTTTAGTTTGAAATGTGTATTAAAGTTACATTTTTGAGTTACAAGAGTCTTATAACTACTTGAA TTTTTAGTTAAAATGTCTTAATGTAGGTTGTAGTCACTTTAGATGGAAAATTACCTCACATCTGTTTTCTTCAGT ATTTATGGTCTAATTTGTATTTAAACATATGCACACATATAAAAGTTGATACTGTGGCAGTAAACTATTAAAAGT TTTCACTGTT

989/2825 FIGURE 872

 ${\tt MPGKKARKNAQPSPARAPAELEVECATQLRRFGDKLNFRQKLLNLISKLFCSGT}$

/

990/2825 FIGURE 873

GCTCCTCACAGGCGCGGGTTGCTTCACCCACACCGTCATGTACTACGGCCACTACAGTAACGCCACGCTGAACCA CTACCTCTCCACTGTGGGCGTGAGCTTCTTTATCACCTGCATCACCCTGGTGTACAGCATCGCTCACTCTTTCGG GGAGAGCTACCGGGTGGGCACCACCTCTGGCATCCACGCCATCACCGTCTTCTGCTCCTGGGACTACAAGGTGAC GCAGAAGCGGGCCTCCCGCCTCCAGCAGGACATATTCGCACCCGGCTGAAGGAGCTGCTGGCCGAGTGGCAGCT GGGGACCGCGCTGGCCTGGCCGTCCACGTCTTCTCGGAGTTCATGATCCAGAGTCCAGAGGCTGCTGG CCAGGAGGCTGTGCTGCTCGTGCCCCTGGTGGTTGGCCTCCTCAACCTGGGGGCCCCCTACCTGTGCCGTGT CCTGGCCGCCTGGAGCCGCATGACTCCCCGGTACTGGAGGTGTACGTGGCCATCTGCAGGAACCTCATCCTCAA GCTGGCCATCCTGGGGACACTGTGCTACCACTGGCTGGGCCGCAGGGTGGGCGTCCTGCAGGGCCAGTGCTGGGA GGATTTTGTGGGCCAGGAGCTGTACCGGTTCCTGGTGATGGACTTCGTCCTCATGTTGCTGGACACGCTTTTTGG GATCATCAAGCTGCTGCTCGTCTTCTATGTCAAGAAGACCAGCCTTCTGGCCAACTGCCAGGCGCCGCCGCCC CTGGCTGGCCTCACACATGAGCACCGTCTTCCTCACGCTGCTCTGCTTCCCCGCCTTCCTGGGCGCCGCTGTCTT CCTCTGCTACGCCGTCTGGCAGGTGAAGCCCTCGAGCACCTGCGGCCCCTTCCGGACCCTGGACACCATGTACGA GGCCGCAGGTGTGGGTGCGCCACCTGGAGGCGGCAGGCCCCAGGGTCTCCTGGCTGCCCTGGGTGCACCGGTA CCTGATGGAAAACACCTTCTTTGTCTTCCTGGTGTCAGCCCTGCTGCTGGCCGTGATCTACCTCAACATCCAGGT GGTGCGGGGCCAGCGCAAGGTCATCTGCCTGCTCAAGGAGCAGATCAGCAATGAGGGTGAGGACAAAATCTTCTT ${\tt TGCGGCACCCCTGCCTGCTCACAGATGAACAGGATGCCTAC}{\tt GGGGGACGGCGATGGGCCTCACGGGCCCGCCCA}$ GCACCCTGAGACCACACTGTTGCCTCCCAGTGACCCTGCTGGGACACCAGGACAAGGAAGACAGTTTCGCCTCTC AAAA

991/2825 FIGURE 874

MAHSFGESYRVGSTSGIHAITVFCSWDYKVTQKRASRLQQDNIRTRLKELLAEWQLRHSPRSVCGRLRQAAVLGL VWLLCLGTALGCAVAVHVFSEFMIQSPEAAGQEAVLLVLPLVVGLLNLGAPYLCRVLAALEPHDSPVLEVYVAIC RNLILKLAILGTLCYHWLGRRVGVLQGQCWEDFVGQELYRFLVMDFVLMLLDTLFGELVWRIISEKKLKRRRKPE FDIARNVLELIYGQTLTWLGVLFSPLLPAVQIIKLLLVFYVKKTSLLANCQAPRRPWLASHMSTVFLTLLCFPAF LGAAVFLCYAVWQVKPSSTCGPFRTLDTMYEAGRVWVRHLEAAGPRVSWLPWVHRYLMENTFFVFLVSALLLAVI YLNIQVVRGQRKVICLLKEQISNEGEDKIFLINKLHSIYERKEREERSRVGTTEEAAAPPALLTDEQDA

992/2825 FIGURE 875

GAGAGCGGAGGGTGCCAGAGGTAGGGGGCCGAGAAACAAAGTTCCCGGGGCTTCCTCCGGGGCCGCGGTCGGGGC TGCGCGTTTGACCGCCCCCCTCCTCGCGAAGCAATCGCTTCCAAACTCCTGCGCGCGGTCATCCTCGGGCCGCCCC GGCTCGGGCAAGGGCACCGTGTGCCAGAGGATCGCCCAGAACTTTGGTCTCCAGCATCTCTCCAGCGCCACTTC TTGCGGGAGAACATCAAGGCCAGCACCGAAGTTGGTGAGATGGCAAAGCAGTATATAGAGAAAAGTCTTTTGGTT CCAGACCATGTGATCACACGCCTAATGATGTCCGAGTTGGAGAACAGGCGTGGACAGCACTGGCTCCTTGATGGT TTTCCTAGGACATTAGGACAAGCCGAAGCCCTGGACAAAATCTGTGAAGTGGATCTAGTGATCAGTTTGAATATT CCATTGAAACACTTAAAGATCGTCTCAGCCGCCGTTGGATTCACCCTCCTAGCGGAAGGGTATATAACCTGGAC TTCAATCCACCTCATGTACATGGTATTGATGACGTCACTGGTGAACCGTTAGTCCAGCAGGAGGATGATAAACCC GAAGCAGTTGCTGCCAGGCTAAGACAGTACAAAGACGTGGCAAAGCCAGTCATTGAATTATACAAGAGCCGAGGA GTGCTCCACCAATTTTCCGGAACGGAGACGAACAAATCTGGCCCTACGTTTACACACTTTTCTCAAACAAGATC TGGTGAACTGATAGGAAAACAAATGAGTAGAAAGAGTTCATGAAGAGGCCCTCCTCTGCCTTTCAAAAGGCTGGT CACCTACACATGTTTAAGGTGTCTCTGCACATGTCTCAAGCCCATCACAAGAAAGCAAGTACAGTGTGGATTTCA AATGGTGTGTAACTTCAGCTCCAGCTGGTTTTTGACAGCTGTTGCTGTGGTAATATTTTTGACATGTGATGGTGA ATAGTGGTTGTTTCTCTGGTCTTGAGTGACTGTGTCCACAGTTCATTTTTTTCCGGTAGGAATAACTCCTTTTCT TTTTGGACATCTGTTTTCACTCTTAGGCTTTTAAACAATAGTTATTGCTTTTATCCCTCTCAGATTCTAATAACT GAGAGCGATGGGGCTATATTGAATCTCTGTATGCACTGAGAACTGAGCTATGAAGAGAATCTTATTAAACTGCTG

993/2825 FIGURE 876

MASKLLRAVILGPPGSGKGTVCQRIAQNFGLQHLSSGHFLRENIKASTEVGEMAKQYIEKSLLVPDHVITRLMMS ELENRRGQHWLLDGFPRTLGQAEALDKICEVDLVISLNIPFETLKDRLSRRWIHPPSGRVYNLDFNPPHVHGIDD VTGEPLVQQEDDKPEAVAARLRQYKDVAKPVIELYKSRGVLHQFSGTETNKIWPYVYTLFSNKITPIQSKEAY

994/2825 FIGURE 877

GAGTTGCCCGTGTGTGCAAGCTGCGGCCAGAGGATCTATGATGGCCAGTACCTCCAGGCCCTGAACGCGGACTGG CACGCAGACTGCTTCAGGTGTTGTGACTGCAGTGCCTCCCTGTCGCACCAGTACTATGAGAAGGATGGGCAGCTC TTCTGCAAGAAGGACTACTGGGCCCGCTATGGCGAGTCCTGCCATGGGTGCTCTGAGCAAATCACCAAGGGACTG GTTATGGTGGCTGGGGAGCTGAAGTACCACCCCGAGTGTTTCATCTGCCTCACGTGTGGGACCTTTATCGGTGAC GGGGACACCTACACGCTGGTGGAGCACTCCAAGCTGTACTGCGGGCACTGCTACTACCAGACTGTGGTGACCCCC GTCATCGAGCAGATCCTGCCTGACTCCCCTGGCTCCCCACCCGCCCACACCGTCACCCTGGTGTCCATCCCAGCC TCATCTCATGGCAAGCGTGGACTTTCAGTCTCCATTGACCCCCGCACGGCCCACCGGGCTGTGGCACCGAGCAC GACCGGATCTTGGAAATCAATGGCACGCCCATCCGAAATGTGCCCCTGGACGAGATTGACCTGCTGATTCAGGAA ACCAGCCGCCTGCTCCAGCTGACCCTCGAGCATGACCCTCACGATACACTGGGCCACGGGCTGGGGCCTGAGACC AGCCCCTGAGCTCTCCGGCTTATACTCCCAGCGGGGAGGCGGCCAGCTCTGCCCGGCAGAAACCTGTCTTCGCA AGGACCTGGGTCGCTCTGAGTCCCTCCGCGTAGTCTGCCGGCCACACCGCATCTTCCGGCCGTCGGACCTCATCC ACGGGGAGGTGCTGGGCAAGGGCTGCTTCGGCCAGGCTATCAAGGTGACACCCGTGAGACAGGTGAGGTGATGG TGATGAAGGAGCTGATCCGGTTCGACGAGGAGACCCAGAGGACGTTCCTCAAGGAGGTGAAGGTCATGCGATGCC TGGAACACCCCAACGTGCTCAAGTTCATCGGGGTGCTCTACAAGGACAAGAGGCTCAACTTCATCACTGAGTACA TCAAGGGCGGCACGCTCCGGGGCATCATCAAGAGCATGGACAGCCAGTACCCATGGAGCCAGAGAGTGAGCTTTG CCAAGGACATCGCATCAGGGATGGCCTACCTCCACTGAACATCATCCACCGAGACCTCAACTCCCACAACT GCCTGGTCCGCGAGAACAAGAATGTGGTGGTGGCTGACTTCGGGCTGGCGCGTCTCATGGTGGACGAGAAGACTC AGCCTGAGGGCCTGCGGAGCCTCAAGAAGCCAGACCGCAAGAAGCCGCTACACCGTGGTGGGCAACCCCTACTGGA TGGCACCTGAGATGATCAACGGCCGCAGCTATGATGAGAAGGTGGATGTGTTCTCCTTTGGGATCGTCCTGTGCG AGATCATCGGGCGGGTGAACGCAGACCCTGACTACCTGCCCCGCACCATGGACTTTGGCCTCAACGTGCGAGGAT TCCTGGACCGCTACTGCCCCCCAAACTGCCCCCGAGCTTCTTCCCCATCACCGTGCGCTGTTGCGATCTGGACC CACTGGGCCCACAGCTGGAGCAGCTGGACAGAGGTTTCTGGGAGACCTACCGGCGCGGGGAGAGCGGACTGCCTG CCCACCCTGAGGTCCCCGACTGAGCCAGGGCCACTCAGCTGCCCCTGTCCCCACCTCTGGAGAATCCACCCCAC TTGCCTTCTCCCACCCGTGGACCGCTTCCCCTGCCTTCTCTCTGCGTGGCCCAGAGCCGGCCCAGCTGCACAC ACACACCATGCTCTCGCCCTGCTGTAACCTCTGTCTTGGCAGGGCTGTCCCCTCTTGCTTCTCCTTGCATGAGCT GGAGGGCCTGTGTGAGTTACGCCCCTTTCCACACGCCGCTGCCCCAGCAACCCTGTTCACGCTCCACCTGTCTGG TCCATAGCTCCCTGGAGGCTGGGCCAGGAGCCTCCGAACCATGCCCCATATAACGCTTGGGTGCGTGGGAG GGCGCACATCAGGGCAGAGGCCAAGTTCCAGGTGTCTGTGTTCCCAGGAACCAAATGGGGAGTCTGGGGCCCGTT TTAAAGCCACTTTAGTGAGAAGCAGGTACCAAGCCTCAGGGTGAAGGGGGTCCCTTGAGGGAGCGTGGAGCTGCG GTGCCCTGGCCGGCGATGGGGAGGAGCCGGCTCCGGCAGTGAGAGGATAGGCACAGTGGACCGGGCAGGTGTCCA CCAGCAGCTCAGCCCCTGCAGTCATCTCAGAGCCCCTTCCCGGGCCTCTCCCCCAAGGCTCCCTGCCCCTCCTCA TGCCCCTCTGTCCTCTGCGTTTTTTCTGTGTAATCTATTTTTTAAGAAGAGTTTGTATTATTTTTCATACGGCT CCTCAGTTGAGCATCTAGGAAGTATTAAAACTGTGAAGCTTTCTCAGTGCACTTTGAACCTGGAAAACAATCCCA CAAACCCAGACGCAGAACAATAAAATAAATTCCGTACTCCCCACCC

995/2825 FIGURE 878

MHHQQRLRANGGRGGTGLGAGSGPVSGGHSPLCAHCRGQLLPHPEEPIVMALGQQWHCDCFRCSVCEGHLHNWYF EREGLLYCREDYYGRFGDACOOCMAVITGPVMVAGEHKFHPECFCCTACGSFIGEGESYALVERSKLYCGQCYGK RSCQPADAKARITTAGKPMHSIRLVEIPKDATPGLRVDGVALDDGCPTVRITDLFCNFFLWLTSMAEQCCGAPYR IDVNLTNLHIGDRILEVNGTPVSDSSVEQIDKLIRSNEKMLQLTVEHDPVQVCRSCSQADIQRAMSASTLILPLS TSASSVEVGRERLYKTPGEQGTKARKLRQATNASTTIPPAAGATAMTQLKEKERCSSLSKLLDEQHQAQQHSAHP QLYDLSRTQSCRVVQKPQRIFRATDLVIGEKLGEGFFGKVFKVTHRQSGEVMVLKELHRADEEAQRNFIKEVAVL RLLDHRHVLKFIGVLYKDKKLHMVTEYVAGGCLKELIHDPAOVLPWPORVRLARDIACGMSYLHSMNIIHRDLNS MNCLVREDRSVIVADFGLARSVDAPRLPSGNMTPGGYGSGANSDAPMSPSGTLRRSKSRQRRQRYTVVGNPYWMA ${\tt PEMMKGLKYDEKLDVFSFGIMLCEIIGRVEADPDFMPRNSDFSLNQQEFREKFCAQCPEPFVKVAFVCCDLNPDM}$ RPCFETLHVWLQRLADDLAADRVPPERLLHEIETFQEWYASSEDALSPTSQRSLNNLDELVKSAVDSEISPVEKE KENMVIKPQDIPKSPHLGKDFSPSGERLRDSMRARRRQRFLGAQEERRNLTPDTESKERALKKALKKCRPFGERG YLVDLRAGAELQLEDVRDLNTYSDVDSSCDTSLNYHDVNNLPAAQEDENTVKPGKEELLEESTNKPSNQESQHHR LAIDDMRTRLNQCRSKFEHLEEASRRNFNQSQHSMKNFFKTPPVALKMFQRLEHEAAALNGGNNCPPPPPRTQRI NQTPIFGRKNPPVAIVGQKLQHAESLEDLASSGVAKQLATPAPKRSKATATTKGGQSSNPPLFLPPSLNISVALN ${\tt SNGNVTTTNTNSSCPPSASDWLPKKHKLTLPLPSAQQQRTSSNHRLPMCNNKGKTLKPLPSRTGSQGIPASNCV}$ SPTRSSRPGSPTKHLAORHTAATAORLTNAAATHQOOHQOOSSKTTRLNILSPEKVHRLGARLTDQKQKMREEAA ATASSVGGAGCAAGTAAGSLNGHRTIGSSGTPNSAVGERRRRAAPSPPVRTHFNTRC

996/2825 FIGURE 879

GCGTGTGAAAGCGCGTTTTACCCGAAGGCGCTCTGTGAGGGCCAGCGGGTCCCCTTCGGCCCTGGAGCCGGGGTC ACACGCTCCCCACCGCGTGCGGTCACGAGACGCCCCCAAGGGAGTATCCTGGTACCCGGAAGCCGCGACTCCTGG CCCTGAGCCCGGGCTTAGCCTTCGGGTCCACGTGGCCGGAGGCGGCAGCTGATTGGACGCGGGCCGCCCCACCCC TGGGGCCAGGCCCGCGGCCGACTTCTCCGTGTCGGTGGAGCGCGCTCTGGCTGCCAAGCCGGGCTTGGACACCT ACAGCCTGGGCGGCGCGCGCGCGCGCGCGCGCGCGCGCCCCACGGGCGTGGCGGCCGCCGCGGGGCTGC ACCGCTACCTGCGCGACTTCTGTGGCTGCCACGTGGCCTGGTCCGGCTCTCAGCTGCCGCCTGCCGCGCCACTGC CAGCCGTGCCGGGGGGCTGACCGAGGCCACGCCCAACAGGTACCGCTATTACCAGAATGTGTGCACGCAAAGCT ACTCCTTCGTGTGGGGACTGGGCCCGCTGGGAGCGAGAGATAGACTGGATGGCGCTGAATGGCATCAACCTGG CACTGGCCTGGAGCGGCCAGGAGGCCATCTGGCAGCGGGTGTACCTGGCCTTGGGCCTGACCCAGGCAGAGATCA ATGAGTTCTTTACTGGTCCTGCCTTCCTGGCCTGGGGGGGCGAATGGGCAACCTGCACACCTGGGATGGCCCCCTGC CCCCTCCTGGCACATCAAGCAGCTTTACCTGCAGCACCGGGTCCTGGACCAGATGCGCTCCTTCGGCATGACCC CAGTGCTGCCTGCATTCGCGGGGCATGTTCCCGAGGCTGTCACCAGGGTGTTCCCTCAGGTCAATGTCACGAAGA TGGGCAGTTGGGGCCACTTTAACTGTTCCTACTCCTGCTCCTTCTTGGCTCCGGAAGACCCCATATTCCCCA TCATCGGGAGCCTCTTCCTGCGAGAGCTGATCAAAGAGTTTGGCACAGACCACATCTATGGGGCCGACACTTTCA ATGAGATGCAGCCACCTTCCTCAGAGCCCTCCTACCTTGCCGCAGCCACCACTGCCGTCTATGAGGCCATGACTG TGTATACCCGCACTGCCTCCTTCCAGGGCCAGCCCTTCATCTGGTGCATGCTGCACAACTTTGGGGGAAACCATG GTCTTTTTGGAGCCCTAGAGGCTGTGAACGGAGGCCCAGAAGCTGCCCGCCTCTTCCCCAACTCCACCATGGTAG GCACGGGCATGGCCCCGAGGGCATCAGCCAGAACGAAGTGGTCTATTCCCTCATGGCTGAGCTGGGCTGGCGAA AGGACCCAGTGCCAGATTTGGCAGCCTGGGTGACCAGCTTTGCCGCCCGGCGGTATGGGGTCTCCCACCCGGACG CAGGGGCAGCGTGGAGGCTACTGCTCCGGAGTGTGTACAACTGCTCCGGGGAGGCCTGCAGGGGCCACAATCGTA GCCCGCTGGTCAGGCGGCCGTCCCTACAGATGAATACCAGCATCTGGTACAACCGATCTGATGTGTTTGAGGCCT GGCGGCTGCTCACATCTGCTCCCTCCCTGGCCACCAGCCCCGCCTTCCGCTACGACCTGCTGGACCTCACTC GGCAGGCAGTGCAGGAGCTGGTCAGCTTGTACTATGAGGAGGCAAGAAGCGCCTACCTGAGCAAGGAGCTGGCCT CCCTGTTGAGGGCTGGAGGCGTCCTGGCCTATGAGCTGCTGCCGGCACTGGACGAGGTGCTGGCTAGTGACAGCC AGAACAGCCGCTACCAGCTGACCTTGTGGGGGCCAGAAGGCAACATCCTGGACTATGCCAACAAGCAGCTGGCGG GGTTGGTGGCCAACTACTACACCCCTCGCTGGCGGCTTTTCCTGGAGGCGCTGGTTGACAGTGTGGCCCAGGGCA ACCCCAGCCAGCCGCGAGGAGACACTGTGGACCTGGCCAAGAAGATCTTCCTCAAATATTACCCCGGCTGGGTGG $\texttt{CCGGCTCTTGG} \underline{\textbf{TGA}} \\ \texttt{TAGATTCGCCACCACTGGGCCTTGTTTTCCGCTAATTCCAGGGCAGATTCCAGGGCCCAGA} \\$ GCTGGACAGACATCACAGGATAACCCAGGCCTGGGAGGAGGCCCCACGGCCTGCTGGTGGGGTCTGACCTGGGGG

997/2825 FIGURE 880

MEAVAVAAAVGVLLLAGAGGAAGDEAREAAAVRALVARLLGPGPAADFSVSVERALAAKPGLDTYSLGGGGAARV RVRGSTGVAAAAGLHRYLRDFCGCHVAWSGSQLRLPRPLPAVPGELTEATPNRYRYYQNVCTQSYSFVWWDWARW EREIDWMALNGINLALAWSGQEAIWQRVYLALGLTQAEINEFFTGPAFLAWGRMGNLHTWDGPLPPSWHIKQLYL QHRVLDQMRSFGMTPVLPAFAGHVPEAVTRVFPQVNVTKMGSWGHFNCSYSCSFLLAPEDPIFPIIGSLFLRELI KEFGTDHIYGADTFNEMQPPSSEPSYLAAATTAVYEAMTAVDTEAVWLLQGWLFQHQPQFWGPAQIRAVLGAVPR GRLLVLDLFAESQPVYTRTASFQGQPFIWCMLHNFGGNHGLFGALEAVNGGPEAARLFPNSTMVGTGMAPEGISQ NEVVYSLMAELGWRKDPVPDLAAWVTSFAARRYGVSHPDAGAAWRLLLRSVYNCSGEACRGHNRSPLVRRPSLQM NTSIWYNRSDVFEAWRLLLTSAPSLATSPAFRYDLLDLTRQAVQELVSLYYEEARSAYLSKELASLLRAGGVLAY ELLPALDEVLASDSRFLLGSWLEQARAAAVSEAEADFYEQNSRYQLTLWGPEGNILDYANKQLAGLVANYYTPRW RLFLEALVDSVAQGIPFQQHQFDKNVFQLEQAFVLSKQRYPSQPRGDTVDLAKKIFLKYYPGWVAGSW

998/2825 FIGURE 881

 ${\tt CCTGGCTGTGGTCGGGGGCAGTGGGCCAGTGGAAGCCCCAGGTGGAAGCAGGCGGAGGACAT}$ TAGAGACATCTACGACTTCCGAGATGTTCTGGGCACGGGGGCCTTCTCGGAGGTGATCCTGGCAGAAGATAAGAG GATTGCTGTCCTGCACAAGATCAAGCACCCCAACATTGTAGCCCTGGATGACATCTATGAGAGTGGGGGCCACCT CTACCTCATCATGCAGCTGGTGTCGGGTGGGGAGCTCTTTGACCGTATTGTGGAAAAAGGCTTCTACACGGAGCG GGACGCCAGCCCCCATCTTCCAGGTGCTGGATGCTGTGAAATACCTGCATGACCTGGGCATTGTACACCGGGA TCTCAAGCCAGAGAATCTGCTGTACTACAGCCTGGATGAAGACTCCAAAATCATGATCTCCGACTTTGGCCTCTC CAAGATGGAGGACCCGGGCAGTGTGCTCCCACCGCCTGTGGAACTCCGGGATACGTGGCCCCTGAAGTCCTGGC $\verb|CCAGAAGCCCTACAGCAAGGCTGTGGATTGCTGGTCCATAGGTGTCATCGCCTACATCTTGCTCTGCGGTTACCC| \\$ TCCCTTCTATGACGAGAATGATGCCAAACTCTTTGAACAGATTTTGAAGGCCGAGTACGAGTTTGACTCTCCTTA $\verb|CTGGGACGACATCTCTGACTCTGCCAAAGATTTCATCCGGCACTTGATGGAGAAGGACCCAGAGAAAAGATTCAC| \\$ CTGTGAGCAGGCCTTGCAGCACCCATGGATTGCAGGAGATACAGCTCTAGATAAGAATATCCACCAGTCGGTGAG ${\tt TGAGCAGATCAAGAAGAACTTTGCCAAGAGCAAGTGGAAGCCATCAATGCCACGGCTGTGGTGCGGCACAT}$ GGCTGGGGGGCCGGCAGCTGCTGCTGCTGCGTGCGTGGAGCCGGGCACAGAACTGTCCCCCACACT $\texttt{GCCCCACCAGCTC} \underline{\textbf{TAG}} \texttt{GGCCCTGGACCTCGGGTCATGATCCTCTGCGTGGGAGGGCTTGGGGGCCAGCCTGCTCCC}$ CTTCCCTCCCTGAACCGGGAGTTTCTCTGCCCTGTCCCCTCACCTGCTTCCCTACCACTCCTCACTGCATTT Α

999/2825 FIGURE 882

MLGAVEGPRWKQAEDIRDIYDFRDVLGTGAFSEVILAEDKRTQKLVAIKCIAKEALEGKEGSMENEIAVLHKIKH PNIVALDDIYESGGHLYLIMQLVSGGELFDRIVEKGFYTERDASRLIFQVLDAVKYLHDLGIVHRDLKPENLLYY SLDEDSKIMISDFGLSKMEDPGSVLSTACGTPGYVAPEVLAQKPYSKAVDCWSIGVIAYILLCGYPPFYDENDAK LFEQILKAEYEFDSPYWDDISDSAKDFIRHLMEKDPEKRFTCEQALQHPWIAGDTALDKNIHQSVSEQIKKNFAK SKWKQAFNATAVVRHMRKLQLGTSQEGQGQTASHGELLTPVAGGPAAGCCCRDCCVEPGTELSPTLPHQL

1000/2825 FIGURE 883

CAGAGGGAGGAAGAÄGCGGCGGCGGCGGCGGCGGCGCTCCTCTTTGCAGAGGGGGGAAACTCTTGGGCTGAGAGCA GGAACAACGCGGTAGGCAAGGCGGGCTGCTGGCTCCCCGGCTCCGGCAGCAGCGGCGGCAGCCCGAGCAGCGGC CAAGAAGTGGAAAGAAATCCTGAAGTTCCCTCACATTAGCCAGTGTGAAGACCTCCGAAGGACCATAGACAGAGA TTACTGCAGTTTATGTGACAAGCAGCCAATCGGGAGGCTGCTTTTCCGGCAGTTTTGTGAAACCAGGCCTGGGCT GGAGTGTTACATTCAGTTCCTGGACTCCGTGGCAGAATATGAAGTTACTCCAGATGAAAAACTGGGAGAGAAAGG GAAGGAAATTATGACCAAGTACCTCACCCCAAAGTCCCCTGTTTTCATAGCCCAAGTTGGCCAAGACCTGGTCTC CCAGACGGAGGAGAAGCTCCTACAGAAGCCGTGCAAAGAACTCTTTTCTGCCTGTGCACAGTCTGTCCACGAGTA CCTGAGGGGAAACCATTCCACGAATATCTGGACAGCATGTTTTTTGACCGCTTTCTCCAGTGGAAGTGGTTGGA AAGGCAACCGGTGACCAAAAACACTTTCAGGCAGTATCGAGTGCTAGGAAAAGGGGGCTTCGGGGAGGTCTGTGC CTGCCAGGTTCGGGCCACGGGTAAAATGTATGCCTGCAAGCGCTTGGAGAAGAAGAGGATCAAAAAGAGGAAAGG CTACGAGACCAAGGATGCACTGTGCTTGGTCCTGACCATCATGAATGGGGGTGACCTGAAGTTCCACATCTACAA CATGGGCAACCCTGGCTTCGAGGAGGAGCGGCCTTGTTTTATGCGGCAGAGATCCTCTGCGGCTTAGAAGACCT CCACCGTGAGAACACCGTCTACCGAGATCTGAAAACCTGAAAACATCCTGTTAGATGATTATGGCCACATTAGGAT CTCAGACCTGGGCTTGGCTGAAGATCCCCGAGGGAGACCTGATCCGCGGCCGGTGGGCACTGTTGGCTACAT GGCCCCGAAGTCCTGAACAACCAGAGGTACGGCCTGAGCCCCGACTACTGGGGCCTTGGCTGCCTCATCTATGA GATGATCGAGGGCCAGTCGCCGTTCCGCGGCCGTAAGGAGAAGGTGAAGCGGGAGGAGGTGGACCGCCGGGTCCT GGAGACGGAGGAGGTGTACTCCCACAAGTTCTCCGAGGAGGCCAAGTCCATCTGCAAGATGCTGCTCACGAAAGA TGCGAAGCAGAGGCTGGGCTGCCAGGAGGAGGGGGCTGCAGAGGTCAAGAGACACCCCTTCTTCAGGAACATGAA CTTCAAGCGCTTAGAAGCCGGGATGTTGGACCCTCCCTTCGTTCCAGACCCCCGCGCTGTGTACTGTAAGGACGT GCTGGACATCGAGCAGTTCTCCACTGTGAAGGGCGTCAATCTGGACCACAGACGACGACGACTTCTACTCCAAGTT CTCCACGGGCTCTGTGTCCATCCCATGGCAAAACGAGATGATAGAAACAGAATGCTTTAAGGAGCTGAACGTGTT TGGACCTAATGGTACCCTCCCGCCAGATCTGAACAGAAACCACCCTCCGGAACCGCCCAAGAAAGGGCTGCTCCA GAGACTCTTCAAGCGGCAGCATCAGAACAATTCCAAGAGTTCGCCCAGCTCCAAGACCAGTTTTAACCACCACAT AAACTCAAACCATGTCAGCTCGAACTCCACCGGAAGCAGCTAGTTTCGGCTCTGGCCTCCAAGTCCACAGTGGAA $\tt CCAGCCCAGACCCTTCTCCTTAGAAGTGGAAGTAGTGGAGCCCCTGCTCTGGTGGGGCTGCCAGGGGAGACCCCG$ GGAGCCGGAAGGAGGCCGTCCATCCCGTCGACGTAGAACCTCGAGGTTTCTCAAAGAAATTTCCACTCAGGTCTG TTTTCCGAGGCGGCCCCGGGCGGGTGGATTGGATTTGTCTTTGGTGAACATTGCAATAGAAATCCAATTGGATAC GACAACTTGCACGTATTTTAATAGCGTCATAACTAGAACTGAATTTTGTCTTTATGATTTTTAAAGAAAAGTTTT GTAAATTTCTCTACTGTCTCAGTTTACATTTTCGGTATATTTGTATTTAAATGAAGTGAGACTTTGAGGGTGTAT AATGTGACTCAAGACTTCCAGAGCCTCAAATGAGAAAATGTCTTTATTAAATGTAGAAAGTGATCCATACTTCAA AAAAAAA

1001/2825 FIGURE 884

MELENIVANTVLLKAREGGGGKRKGKSKKWKEILKFPHISQCEDLRRTIDRDYCSLCDKQPIGRLLFRQFCETRP GLECYIQFLDSVAEYEVTPDEKLGEKGKEIMTKYLTPKSPVFIAQVGQDLVSQTEEKLLQKPCKELFSACAQSVH EYLRGEPFHEYLDSMFFDRFLQWKWLERQPVTKNTFRQYRVLGKGGFGEVCACQVRATGKMYACKRLEKKRIKKR KGESMALNEKQILEKVNSQFVVNLAYAYETKDALCLVLTIMNGGDLKFHIYNMGNPGFEEERALFYAAEILCGLE DLHRENTVYRDLKPENILLDDYGHIRISDLGLAVKIPEGDLIRGRVGTVGYMAPEVLNNQRYGLSPDYWGLGCLI YEMIEGQSPFRGRKEKVKREEVDRRVLETEEVYSHKFSEEAKSICKMLLTKDAKQRLGCQEEGAAEVKRHPFFRN MNFKRLEAGMLDPPFVPDPRAVYCKDVLDIEQFSTVKGVNLDHTDDDFYSKFSTGSVSIPWQNEMIETECFKELN VFGPNGTLPPDLNRNHPPEPPKKGLLQRLFKRQHQNNSKSSPSSKTSFNHHINSNHVSSNSTGSS

1002/2825 FIGURE 885

CTCCTTATTCTCAAAAGGGGGAGCTGGGGAGCCTCCCCCTACCCTGGGGCCTCCCCTGCCCCTGCCCTGCC TGGCCGTCACCACTCCCCAGAGGGCACAGGGCTCTGCTGTGCCTCAGAGCAAAAGTCCCAGAGCCAGCAGAGCAG AGCCCCAAGACCCCGGGGGCCATGGGCGGGGATCTGGTGCTTGGCCTGGGGGGCCTTGAGACGCCCAAAGCCGCTTG CTGGAGCAGGAGAGTCTCTGGCCGGCTGGGCACTGGTGCTGGCAGGAACTGGCATTGGACTCATGGTGCTGCAT GCAGAGATGCTGTGGTCGGGGGGGTGCTCGTGGGCGCTCTACCTGTTCCTGGTTAAATGCACGATCAGCATTTCC ACCTTCTTACTCCTCTGCCTCATCGTGGCCTTTCATGCCAAAGAGGTCCAGCTGTTCATGACCGACAACGGGCTG CCGGCGCCCGTGCGGGCCCGTGCGTGCAGGATTTAGGGGCGCCGCTGACCTCCCCGCAGCCCTGGCCGGGA TTCCTGGGCCAAGGGGAAGCGCTGCTGTCCCTGGCCATGCTGCTGCTCTACCTGGTGCCCCGCGCCGTGCTC TTCGTGGCCAAGCTTTACATGAACACGCACCCTGGCCGCCTGCTGGTCGGCCTCACGCTTGGCCTCTGGCTGACC ACCGCCTGGGTGCTGTCCGTGGCCGAGAGGCAGGCTGTTAATGCCACTGGGCACCTTTCAGACACACTTTGGCTG ATCCCCATCACATTCCTGACCATCGGCTATGGTGACGTGGTGCCGGGCACCATGTGGGGCAAGATCGTCTGCCTG TGCACTGGAGTCATGGGTGTCTGCTGCACAGCCCTGCTGGTGGCCGGGTGGCCCGGAAGCTGGAGTTTAACAAG GCAGAGAAGCACGTGCACAACTTCATGATGGATATCCAGTATACCAAAGAGATGAAGGAGTCCGCTGCCCGAGTG CTGCTGGCCGCCATCAACGCGTTCCGCCAGGTGCGGCTGAAACACCGGAAGCTCCGGGAACAAGTGAACTCCATG GTGGACATCTCCAAGATGCACATGATCCTGTATGACCTGCAGCAGAATCTGAGCAGCTCACACCGGGCCCTGGAG $\tt CTTCCAGAACCCAGCCAGCAGTCCAAG{\color{red}{TAG}} CTGGACCCACGAGGAGCAACCAGGCTACTTTCCCCAGTACTGAGG$ TGGTGGACATCGTCTCTGCCACTCCTGACCCAGCCCTGAACAAGCACCTCAAGTGCAAGGACCAAAGGGGGCCC TGGCTTGGAGTGGGTTGGCTGATGGCTGCTGGAGGGGACGCTGGCTAAAGTGGGTAGGCCTTGGCCCACCT GAGGCCCCAGGTGGGAACATGGTCACCCCCACTCTGCATACCCTCATCAAAAACACTCTCACTATGCTGCTATGG ACGACCTCCAGCTCTCAGTTACAAGTGCAGGCGACTGGAGGCAGGACTCCTGGGTCCCTGGGAAAGAGGGTACTA GGGGCCCGGATCCAGGATTCTGGGAGGCTTCAGTTACCGCTGGCCGAGCTGAAGAACTGGGTATGAGGCTGGGGC GGGGCTGGAGGTGGCCCCCCTGGTGGGACAACAAGAGGACACCATTTTTCCAGAGCTGCAGAGAGCACCTGGT GGGGAGGAAGAAGTGTAACTCACCAGCCTCTGCTCTTATCTTTGTAATAAATGTTAAAGCCAG

1003/2825 FIGURE 886

MGGDLVLGLGALRRRKRLLEQEKSLAGWALVLAGTGIGLMVLHAEMLWFGGCSWALYLFLVKCTISISTFLLLCL IVAFHAKEVQLFMTDNGLRDWRVALTGRQAAQIVLELVVCGLHPAPVRGPPCVQDLGAPLTSPQPWPGFLGQGEA LLSLAMLLRLYLVPRAVLLRSGVLLNASYRSIGALNQVRFRHWFVAKLYMNTHPGRLLLGLTLGLWLTTAWVLSV AERQAVNATGHLSDTLWLIPITFLTIGYGDVVPGTMWGKIVCLCTGVMGVCCTALLVAVVARKLEFNKAEKHVHN FMMDIQYTKEMKESAARVLQEAWMFYKHTRRKESHAARRHQRKLLAAINAFRQVRLKHRKLREQVNSMVDISKMH MILYDLQQNLSSSHRALEKQIDTLAGKLDALTELLSTALGPRQLPEPSQQSK

1004/2825 FIGURE 887

1005/2825 FIGURE 888

 ${\tt MRLFLSLPVLVVVLSIVLEGPAPAQGTPDVSSALDKLKEFGNTLEDKARELISRIKQSELSAKMREWFSETFQKVKEKLKIDS}$

, ...

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1006/2825 FIGURE 889

GAATACTCCCGGGGCCTGGAAAAGCTGGCCGAGCGCTTCTCCAGCCGTGGAGGCCGCCTGGGGAGCAGCCGGGAG CACCAAAGCTTCCGGAAGGAGCCGTCCCTCCTGTCGCCCTTGCACTGCTGGGCGGTGCTGCAGCACACGCGG CAGCAGAGCCGGGAGAGCGCGCCCTGAGTGAGGTGCTGGCCGGGCCCCTGGCCCAGCGCCTGAGTCACATTGCA GAGGACGTGGGGCCCTGGTCAAGAAGAGCAGGGATCTGGAGCAGCAGCTGCAGGATGAGCTCCTGGAGGTGGTC TCAGAGCTCCAGACGGCCAAGAAGACGTACCAGGCATATCACATGGAGAGCGTGAATGCCGAGGCCAAGCTCCGG GAGGCCGAGCGGCAGGAGAAGCGGGCAGGCCGGAGTGTCCCCACCACCACCGCTGGTGCCACTGAGGCAGGG CCCCTCCGCAAGAGCTCCCTCAAGAAGGGAGGGAGGCTGGTGGAGAAGCGGCCAAGTTCATGGAGCACAAA CTCAAGTGCACAAAGGCGCGCAACGAGTACCTGCTTAGCCTGGCTAGTGTCAACGCTGCTGTCAGTAACTACTAC CTGCATGACGTCTTGGACCTCATGGACTGCTGTGACACAGGGTTCCACCTGGCCCTGGGGCAGGTGCTCCGGAGC TACACGGCCGCTGAGAGCCGCACCCAAGCCTCCCAAGTGCAGGGCCTGGGCAGCCTGGAAGAAGCTGTGGAGGCC $\verb|CTGGATCCTCCAGGGGACAAAGCCAAGGTTCTCGAGGTGCATGCTACCGTCTTCTGTCCCCCGCTGCGCTTTGAC| \\$ TACCACCCCATGATGGGGATGAGGTGGCTGAGATCTGCGTTGAAATGGAGCTGCGGGACGAGATTCTGCCCAGA GAAACCTTCTACCTCACGAAGCTCCAGGAGTATCTGAGTGGACGGAGCATCCTCGCCAAGCTGCAGGCCAAGCAC GAGAAGCTGCAGGAGGCCCTTCAGCGAGGTGACAAGGAGGAGCAGGAGGTGTCTTGGACCCAGTACACACAGAGA AAATTCCAGAAGAGCCGCCCGCCCCAGCTCCCAGTATAACCAGAGACTCTTTGGGGGGAGACATGGAGAAG TTTATCCAGAGCTCAGGCCAGCCTGTGCCCCTGGTGGTGGAGAGCTGCATTCGCTTCATCAACCTCAATGGCCTG CAGCATGAAGGCATCTTCCGGGTATCGGGTGCCCAGCTCCGGGTCTCAGAGATCCGTGATGCCTTCGAGAGAGGG GAGGACCCACTGGTGGAGGGCTGCACTGCCCATGACCTGGACTCGGTGGCCGGGGTGCTGAAGCTCTACTTCCGG AGCCTGGAGCCCCACTCTTCCCCCCAGACCTGTTCGGCGAGCTGCTGGCTTCTTCGGAGCTGGAGGCCACAGCG GAGAGGGTGGAGCACGTGAGCCGCCTGCTGTGGCGGCTGCCCGCGCGGTGCTGGTGGTTCTGCGCTACCTCTTC ACCTTCCTCAACCACCTGGCCCAGTACAGCGATGAGAACATGATGGACCCCTACAACCTGGCCGTGTGCTTCGGG CCCACGCTGCTACCGGTGCCCGCTGGGCAGGACCCGGTGGCGCTGCAGGGCCGGGTGAACCAGCTGGTGCAGACG CTCATAGTGCAGCCCGATCGGGTCTTCCCGCCCCTGACCTCGCCTGGCCCCGTCTACGAGAAGTGCATGGCA CCGCCTTCCGCCAGCTGCCTGGGGGACGCCCAGCTGGAGAGCCTGGGGGGCGGACAATGAGCCGGAAGCTGGAAGCC GAGATGCCCGCACAGGAGGATGACCTGGAGGGGGTCGTGGAGGCTGTGGCCTGCTTTGCCTACACGGGCCGCACA GCCCAGGAGCTGAGCTTCCGGCGGGGGGACGTACTGCGGCTGCACGAGAGGGCCTCGAGCGACTGGTGGCGGGG GAGCACAACGGCATGCGGGGCCTCATCCCCCACAAGTATATCACGCTGCCCGCGGGACGGAGAAGCAGGTGGTG GGCGCAGGGCTGCAGACTGCAGGGGAGTCTGGGAGCAGTCCCGAGGGCCTCCTGGCATCGGAGCTGGTCCACCGG CCAGAGCCATGCACCTCACCTGAGGCCATGGGACCCTCTGGACACAGACGACGCTGCTTGGTCCCAGCCTCCCCA GAGCAACACGTGGAGGTGGATAAGGCTGTGGCACAGAACATGGACTCTGTGTTTAAGGAGCTCTTGGGAAAGACC TCTGTCCGCCAGGGCCTTGGGCCAGCATCTACCACCTCTCCCAGTCCTGGGCCCCGAAGCCCAAAGGCCCCGCCC AGCAGCCGCCTGGGCAGGAACAAGGCTTCTCCCGGGGCCCTGGGGCCCCAGCCTCACCCTCAGCTTCCCACCCC $\texttt{CAGGGCCTAGACACGACCCCAAGCCACAC} \underline{\textbf{TGA}} \texttt{GGTGCCGCTGCTGGAGATGCGTGCCCCGGCGGCTACCCGCT}$ GGACCGGCCACTCTCCCCAGCCCCTTGCTTCTCTCCAGCCCTGTCCAGCAAGTGCAGGGTGCCTGCACTTCACC CCAGGCCCCTTTTGGAGGCAGCTGAGGAAGGATGCTGGGGAAAGCCCTCTTCTGCAGCTTTGTGGAAGGCTGATC AGTGGCTGCTGGGTGGCGGGTACCCTTGCTCAGATGCCTGGCAGGGCTGGGTGGCGATTCATAAAGACCTCGTGT TGATTCCCCG

1007/2825 FIGURE 890

MAAHGKLRRERGLQAEYETQVKEMRWQLSEQLRCLELQGELRRELLQELAEFMRRRAEVELEYSRGLEKLAERFS
SRGGRLGSSREHQSFRKEPSLLSPLHCWAVLLQHTRQQSRESAALSEVLAGPLAQRLSHIAEDVGRLVKKSRDLE
QQLQDELLEVVSELQTAKKTYQAYHMESVNAEAKLREAERQEEKRAGRSVPTTTAGATEAGPLRKSSLKKGGRLV
EKRQAKFMEHKLKCTKARNEYLLSLASVNAAVSNYYLHDVLDLMDCCDTGFHLALGQVLRSYTAAESRTQASQVQ
GLGSLEEAVEALDPPGDKAKVLEVHATVFCPPLRFDYHPHDGDEVAEICVEMELRDEILPRAQNIQSRLDRQTIE
TEEVNKTLKATLQALLEVVASDDGDVLDSFQTSPSTESLKSTSSDPGSRQAGRRRGQQQETETFYLTKLQEYLSG
RSILAKLQAKHEKLQEALQRGDKEEQEVSWTQYTQRKFQKSRQPRPSSQYNQRLFGGDMEKFIQSSGQPVPLVVE
SCIRFINLNGLQHEGIFRVSGAQLRVSEIRDAFERGEDPLVEGCTAHDLDSVAGVLKLYFRSLEPPLFPPDLFGE
LLASSELEATAERVEHVSRLLWRLPAPVLVVLRYLFTFLNHLAQYSDENMMDPYNLAVCFGPTLLPVPAGQDPVA
LQGRVNQLVQTLIVQPDRVFPPLTSLPGPVYEKCMAPPSASCLGDAQLESLGADNEPELEAEMPAQEDDLEGVVE
AVACFAYTGRTAQELSFRRGDVLRLHERASSDWWRGEHNGMRGLIPHKYITLPAGTEKQVVGAGLQTAGESGSSP
EGLLASELVHRPEPCTSPEAMGPSGHRRRCLVPASPEQHVEVDKAVAQNMDSVFKELLGKTSVRQGLGPASTTSP
SPGPRSPKAPPSSRLGRNKGFSRGPGAPASPSASHPQGLDTTPKPH

1008/2825 FIGURE 891A

GGGAACTTGGATTAGGTGGAGAGGCAGTTGGGGGGCCTCGTTGTTTTGCGTCTTAGTTCCGCCCTCCTGTCCATC $\tt AGGAGAAGGAAAGGATAAACCCTGGGCC{\color{red} \underline{\textbf{ATG}}} \\ AGGCTACCCTGCTCCTGGTTTTTTGCCTCTGTCATTCCGGGTGC$ TGTTCTCCTACTGGACACCAGGCAATTTTTAATCTATAATGAAGATCACAAGCGCTGCGTGGATGCAGTGAGTCC CAGTGCCGTCCAAACCGCAGCTTGCAACCAGGATGCCGAATCACAGAAATTCCGATGGGTGTCCGAATCTCAGAT TATGAGTGTTGCATTTAAATTATGCCTGGGAGTGCCATCAAAAACAGACTGGGTTGCTATCACTCTCTATGCCTG TGACTCAAAAAGTGAATTTCAGAAATGGGAGTGCAAAAATGACACACTTTTGGGGATCAAAGGAGAAGATTTATT TTTTAACTACGGCAACAGACAAGAAAAGAATATTATGCTCTACAAGGGATCGGGTTTATGGAGCAGGTGGAAGAT CTATGGAACCACAGACAATCTGTGCTCCAGAGGTTATGAAGCCATGTATACGCTACTAGGCAATGCCAATGGAGC GCTCTGGTGCGGAACCACTACTGACTATGACACAGACAAGCTATTTGGATATTGTCCATTGAAATTTGAGGGCAG TGAAAGCTTATGGAATAAAGACCCGCTGACCAGCGTTTCCTACCAGATAAACTCCAAATCCGCTTTAACGTGGCA CCAAGCGAGGAAAAGCTGCCAACAACAGAACGCTGAGCTCCTGAGCATCACAGAGATACATGAGCAAACATACCT GACAGGATTAACCAGTTCCTTGACCTCAGGACTCTGGATTGGACTTAACAGTCTGAGCTTCAACAGCGGTTGGCA GTGGAGTGACCGCAGTCCTTTCCGATATTTGAACTGGTTACCAGGAAGTCCATCAGCTGAACCTGGAAAAAGCTG TGTGTCACTAAATCCTGGAAAAAATGCTAAATGGGAAAATCTGGAATGTGTTCAGAAACTGGGCTATATTTGCAA AAAGGGCAACACCACTTTAAATTCTTTTGTTATTCCCTCAGAAAGTGATGTGCCTACTCACTGTCCTAGTCAGTG CAGGAAGGAAGGCGGTGACCTCACAAGTATCCACACCATCGAGGAATTGGACTTTATTATCTCCCAGCTAGGATA TGAGCCAAATGACGAATTGTGGATCGGCTTAAATGACATTAAGATTCAAATGTACTTTGAGTGGAGTGATGGGAC GAAAGGCAAGGATGGGTACTGGGCAGATCGGGGCTGTGAGTGGCCTCTTGGCTACATCTGCAAGATGAAATCACG AAGCCAAGGTCCAGAAATAGTGGAAGTCGAAAAAGGCTGCAGGAAAGGCTGGAAAAAACATCACTTTTACTGCTA TATGATTGGACATACGCTTTCAACATTTGCAGAAGCAAACCAAACCTGTAATAATGAGAATGCTTATTTAACAAC TATTGAAGACAGATATGAACAAGCCTTCCTGACTAGTTTCGTTGGCTTAAGGCCTGAAAAATATTTCTGGACAGG ACTTTCAGATATACAAACCAAAGGGACTTTTCAGTGGACCATCGAGGAAGAGGTTCGGTTCACCCACTGGAATTC ${\tt AGATATGCCAGGGCGAAAGCCAGGGTGTTTTGCCATGAGAACCGGGATTGCAGGGGGCTTATGGGATGTTTTGAA}$ TCCCGAACCCAAATGTCCGGAGGATTGGGGCGCCAGCAGTAGAACAAGCTTGTGTTTCAAGCTGTATGCAAAAGG ATTGACATATGGAAGCCCTTCAGAAGGTTTTACTTGGAGTGATGGTTCTCCTGTTTCATATGAAAACTGGGCTTA TGGAGAACCTAATAATTATCAAAATGTTGAATACTGTGGTGAGCTGAAAGGTGACCCTACTATGTCTTGGAATGA TATTAATTGTGAACACCTTAACAACTGGATTTGCCAGATACAAAAAGGACAAAACACCAAAAACCTGAGCCAACACC AGAGAAGGAAACCATGGACAATGCGCGAGCGTTTTGCAAGAGGAATTTTGGTGATCTTGTTTCTATTCAAAGTGA AAGTGAAAAGAAGTTTCTATGGAAATATGTAAACAGAAATGATGCACAGTCTGCATATTTTATTGGTTTATTGAT CAGCTTGGATAAAAAGTTTGCTTGGATGGATGGAAGCAAAGTGGATTACGTGTCTTGGGCCACAGGTGAACCCAA TTTTGCAAATGAAGATGAAAACTGTGTGACCATGTATTCAAATTCAGGGTTTTGGAATGACATTAACTGTGGCTA TCCAAACGCCTTCATTTGCCAGCGACATAACAGTAGTATCAATGCTACCACAGTTATGCCTACCATGCCCTCGGT CCCATCAGGGTGCAAGGAAGGTTGGAATTTCTACAGCAACAAGTGTTTCAAAATCTTTGGATTTATGGAAGAAGA AAGAAAAATTGGCAAGAGGCACGAAAAGCTTGTATAGGCTTTGGAGGGAATCTGGTCTCCATACAAAATGAAAA AGAGCAAGCATTTCTTACCTATCACATGAAGGACTCCACTTTCAGTGCCTGGACTGGGCTGAATGATGTCAATTC AGAACACACGTTCCTTTGGACGGATGGACGAGGAGTCCATTACACAAACTGGGGGAAAGGTTACCCTGGTGGAAG AAGAAGCAGTCTTTCTTATGAAGATGCTGACTGTTGTTATTATTGGAGGTGCATCAAATGAAGCAGGAAAATG AGCAACGATTCAAACAGATGGCTTTGTTAAATATGGCAAAAGCAGCTATTCACTCATGAGACAAAAATTTCAATG GCATGAAGCGGAGACATACTGCAAGCTTCACAATTCCCTTATAGCCAGCATTCTGGATCCCTACAGTAATGCATT TGCGTGGCTGCAGATGGAAACATCTAATGAACGTGTGTGGATCGCCCTGAACAGTAACTTGACTGATAATCAATA

1009/2825 FIGURE 891B

CACTTGGACTGATAAGTGGAGGGTGAGGTACACTAACTGGGCTGCTGATGAGCCCAAATTGAAATCAGCATGTGT TTATCTGGATCTTGATGGCTACTGGAAGACAGCACATTGCAATGAAAGTTTTTACTTTCTCTGTAAAAGATCAGA TGAAATCCCTGCTACTGAACCCCCACAACTGCCTGGCAGATGCCCGGAGTCAGATCACACAGCATGGATTCCTTT CCATGGTCACTGTTACTATATTGAGTCCTCATATACAAGAAACTGGGGCCCAAGCTTCTCTGGAATGTCTTCGAAT GGGTTCCTCTCTGGTTTCCATTGAAAGTGCTGCAGAATCCAGTTTTCTGTCATATCGGGTTGAGCCACTTAAAAG $\tt CTTTGTCAACTGGAACACAGGAGATCCCTCTGGTGAACGGAATGATTGTGTAGCTTTACATGCGTCTTCTGGGTT$ TTGGAGTAATATTCACTGTTCTTCCTACAAAGGATATATTTGTAAAAGACCAAAAATTATTGATGCTAAACCTAC TCATGAATTACTTACAACAAAAGCTGACACAAGGAAGATGGACCCTTCTAAACCGTCTTCCAACGTGGCCGGAGT AGTCATCATTGTGATCCTCCTGATTTTAACGGGTGCTGGCCTTGCCGCCTATTTCTTTTATAAGAAAAGACGTGT GAAAGATCTCGTGGGCAATATTGAACAGAATGAACACTCGGTCATC**TAG**TACCTCAATGCGATTCTGAGATATTT GAATTTCATAAAATTGTAACTGAAATTTAAAATTTTTAGTTCAATGTGATTGTTTTCTTTAAAATGAGTACTGAA TTGTACTGGTCTGTCCTTTTTCCTTTGCCTAATTGAAGAAATAATTGCTTGTTTTCTAGCCTGGCAAGATATTT TCATAAAAGAGGGATAACAATGCTGATTACTACCTTTTAAAATATTTTAGATAAATGCACACCACCACCACCACCA CATCTAAGCATTAGTGATGGGTAGCTGATGTCAGCTTCATGTGGATTTTAAGCACTCTAGAAACAATGAAGCTTC TTGGCATATTTTAAGGAGCTCCCAAAATGTGTTACCTATTAAATTGTAACTCAGCAAGTAGAAGACCATTTGAAA AGTCAGGTACAAATTTCCTCAAGTGGCATAAAAATGTAGTCAGTTTTCTCTTTTACCAGTTTTTATTTCCACTCC AATTATTTAGAACTTTATTTGTACATGTGCAGAAGAATAAGGCAGCTGAGAATCTTGTTTCCCCCAAGAGAGTTT TACAGGCTGAGTGTTGCAAATGTGTTCTTTGTCCTGTTATATGTATATCAGGAATACAAGGATGTGAAATAAAAC TGTAAATTTGCATAACTGGATGTACTTAGATAATGTGAAATAAACATTAAAGACAAGGTCTATTTTTAATAAAAA AAAAAAAAA

1010/2825 FIGURE 892

MRLPLLLVFASVIPGAVLLLDTRQFLIYNEDHKRCVDAVSPSAVQTAACNQDAESQKFRWVSESQIMSVAFKLCL GVPSKTDWVAITLYACDSKSEFQKWECKNDTLLGIKGEDLFFNYGNRQEKNIMLYKGSGLWSRWKIYGTTDNLCS RGYEAMYTLLGNANGATCAFPFKFENKWYADCTSAGRSDGWLWCGTTTDYDTDKLFGYCPLKFEGSESLWNKDPL TSVSYQINSKSALTWHQARKSCQQQNAELLSITEIHEQTYLTGLTSSLTSGLWIGLNSLSFNSGWQWSDRSPFRY LNWLPGSPSAEPGKSCVSLNPGKNAKWENLECVQKLGYICKKGNTTLNSFVIPSESDVPTHCPSQWWPYAGHCYK IHRDEKKIQRDALTTCRKEGGDLTSIHTIEELDFIISQLGYEPNDELWIGLNDIKIQMYFEWSDGTPVTFTKWLR GEPSHENNRQEDCVVMKGKDGYWADRGCEWPLGYICKMKSRSQGPEIVEVEKGCRKGWKKHHFYCYMIGHTLSTF ${ t AEANQTCNNENAYLTTIEDRYEQAFLTSFVGLRPEKYFWTGLSDIQTKGTFQWTIEEEVRFTHWNSDMPGRKPGC}$ VAMRTGIAGGLWDVLKCDEKAKFVCKHWAEGVTHPPKPTTTPEPKCPEDWGASSRTSLCFKLYAKGKHEKKTWFE SRDFCRALGGDLASINNKEEQQTIWRLITASGSYHKLFWLGLTYGSPSEGFTWSDGSPVSYENWAYGEPNNYQNV EYCGELKGDPTMSWNDINCEHLNNWICQIQKGQTPKPEPTPAPQDNPPVTEDGWVIYKDYQYYFSKEKETMDNAR AFCKRNFGDLVSIQSESEKKFLWKYVNRNDAQSAYFIGLLISLDKKFAWMDGSKVDYVSWATGEPNFANEDENCV TMYSNSGFWNDINCGYPNAFICQRHNSSINATTVMPTMPSVPSGCKEGWNFYSNKCFKIFGFMEEERKNWQEARK ACIGFGGNLVSIQNEKEQAFLTYHMKDSTFSAWTGLNDVNSEHTFLWTDGRGVHYTNWGKGYPGGRRSSLSYEDA DCVVIIGGASNEAGKWMDDTCDSKRGYICQTRSDPSLTNPPATIQTDGFVKYGKSSYSLMRQKFQWHEAETYCKL HNSLIASILDPYSNAFAWLQMETSNERVWIALNSNLTDNQYTWTDKWRVRYTNWAADEPKLKSACVYLDLDGYWK TAHCNESFYFLCKRSDEIPATEPPQLPGRCPESDHTAWIPFHGHCYYIESSYTRNWGQASLECLRMGSSLVSIES AAESSFLSYRVEPLKSKTNFWIGLFRNVEGTWLWINNSPVSFVNWNTGDPSGERNDCVALHASSGFWSNIHCSSY KGYICKRPKIIDAKPTHELLTTKADTRKMDPSKPSSNVAGVVIIVILLILTGAGLAAYFFYKKRRVHLPQEGAFE NTLYFNSQSSPGTSDMKDLVGNIEQNEHSVI

1011/2825 FIGURE 893

AGCGCAGCACTCCCCGCTCGTTGGCCCGGGTATCCCAGCGCGGACCCACGCGATACGCTGACGCCCGACGCCGA TCCGGCCGAGCCAAGTAAGGGGGACGCCCGAGACGGAGAAGGGAGAGTGGGAGTTTCCCAGCCCGCAGAACT $\verb|TTCGAAGTTGAGAAGAGAACCCCTGGAACGTGCGCTCAGCACTGGGATTTTCTGGACTCAACG| \textbf{ATC} \\ \texttt{ACTCTGAAT}$ AATGTCACCATGCGCCAGGGCACTGTGGGCATGCAGCCACAGCAGCGCGCTGGAGCATCCCAGCTGATGGCAGG CATCTGATGGTCCAGAAAGAGCCCCACCAGTACAGCCACCGCCAACCGCCATTCTGCTACCCCTGAGGACCACTGC CGCCGAAGCTGGTCCTCTGACTCCACAGACTCAGTCATCTCCTCTGAGTCAGGGAACACCTACTACCGAGTGGTG CTCATAGGGGAGCAGGGGGGGGCAAGTCCACTCTGGCCAACATCTTTGCAGGTGTGCATGACAGCATGGACAGC GACTGCGAGGTGCTGGGAAAGATACATATGAACGAACCCTGATGGTTGATGGGGAAAGTGCAACGATTATACTC ATTGTCTACTCAATCACAGACCGAGCGAGCTTCGAGAAGGCATCTGAGCTGCGAATCCAGCTCCGCAGGGCCCGG CAGACAGAGGACATTCCCATAATTTTGGTTGGCAACAAAAGTGACTTAGTGCGGTGCCGAGAAGTGTCTGTATCA GAAGGGAGACCTGTGCAGTGGTGTTTGACTGCAAGTTCATCGAGACCTCTGCAGCTGTCCAGCACAACGTGAAG GAGCTGTTTGAGGGCATTGTGCGACAGGTGCGCCTTCGGCGGGACAGCAAGGAGAAGAATGAACGGCGGCTGGCC TACCAGAAAAGGAAGGAGAGCATGCCCAGGAAAGCCAGGCGCTTCTGGGGCAAGATCGTGGCCAAAAACAACAAG TTTCCCCCATTGTAGCAGGGAGCTAGCGTATTAGCCTTGTGGGCAACATGATGCATGGGAAATGAAAGATTTTTG TAAAAAGTCAGTATTTATTTCCAGGAAAAGCCTGACCTTGCTATTTGAACACCCAAGACTCTTTAGAGGGATGTGT TTGGTGTTCACATGTGTTTCTTCTATTTTGGATAGTAGGGAAGTAAAGCTTACAAAGAATGCCTAGAACAAGAAC TTTTCATCATTAAAAATTTTTCCCAGTGTTCTGATATGTGACTTTGAGGCCAATGAGTCATAAACAAATATAAGA AAGCTGTCAATGAGTTTCTTCAAAGGAGGGAAAACTTTCTACGAATCTAAGATCCATGGAGCTAGAATTGTAGAA AAAGGCTTAGCAAACAATTTTTGTTCAATGCCCACCGAGACATATAGAATTGGGAACTGATACATGTGTCCCTTA TAGGCTCAAAAATTATATCTTACAATTTCTTATTTAGGGGGGAAATTATTTGAATCAGATTCTATTTAGTCAAACC ACCTTTTATGTTTTATTTTTTGAATTCATGGAGCCATCATAAAAATATTTTTAAAATCAGAATTATTGATACC CTGTAGTGCAAAATGTCAATTTTTAATGTATAATCAGAAGTCTGAATTTTCATAAAAACATATAGCATAAAAACCT ATTTGATAAAGCATATTTATGAACATATTTCTTAATAAGAAAAATATCCATTTTATTACCATTTTCTATCTTTTT CAAAATATGCAAGTTTTTACCTATATGTCTTATAATAAAAGAAATAAAATATTTGA

1012/2825 FIGURE 894

MTLNNVTMRQGTVGMQPQQQRWSIPADGRHLMVQKEPHQYSHRNRHSATPEDHCRRSWSSDSTDSVISSESGNTY YRVVLIGEQGVGKSTLANIFAGVHDSMDSDCEVLGEDTYERTLMVDGESATIILLDMWENKGENEWLHDHCMQVG DAYLIVYSITDRASFEKASELRIQLRRARQTEDIPIILVGNKSDLVRCREVSVSEGRACAVVFDCKFIETSAAVQ HNVKELFEGIVRQVRLRRDSKEKNERRLAYQKRKESMPRKARRFWGKIVAKNNKNMAFKLKSKSCHDLSVL

1013/2825 FIGURE 895

1014/2825 FIGURE 896

 ${\tt MDCEVNNGSSLRDECITNLLVFGFLQSCSDNSFRRELDALGHELPVLAPQWEGYDELQTDGNRSSHSRLGRIEADS SESQEDIIRNIARHLAQVGDSMDRSIPPGLVNGLALQLRNTSRSEEDRNRDLATALEQLLQAYPRDMEKEKTMLVLALLLAKKVASHTPSLLRDVFHTTVNFINQNLRTYVRSLARNGMD}$

1015/2825 FIGURE 897

GGCACGAGGGACAAAGGCTGCAGGGGGACCCTGCCTGCCCAGGAGGGCCCCTCAGGGAGTAAACCCCTAAGGGGC TCACTTAAGGAGGAGGTAGCCCTGGATTTGAGTGTGAGGAAGCCCACAGCAGAGGCCTCCCCTGTCAAGGCTTCC CGTTCTGTGGAGCATGCCAAGCCTACTGCAGCCATGGATGTGCCAGATGTGGGCAACATGGTGTCAGATCTGCCA GGCCTGAAAAAGATAGACACAGAAGCACCAGGCTTGCCTGGGGTGCCAGTGACCACAGATGCCATGCCAAGGACC CGGATTCTCGCTCAACAGCCCTTGTCTGTGACCTGCTTCAGCCTGGCACTGCCCAGCCCTCCAGCCGTAGCTGTG GCCTCCCTGCCCCTGCTCCAGCTCCATCCCCTGCTCCGGCTCAGGCTCAGCTCCAGCTTCAGCCCGGGATCCA GCTCCAGCTCCAGCTCCAGTTGCAGCCCTGCTCCAGCATCTACTTCAGCCCCAGGGGACTCCCTGGAGCAGCAT TTTACAGGACTACATGCGTCCCTGTGTGATGCTATTTCTGGCTCCGTCGCCCACTCTCCTCCAGAGAAGCTTCGC GAGTGGCTAGAGACGGCTGGGCCCTGGGGCCAGGCTGCGTGGCAGGACTGCCAGGGTGTGCAGGGGCTGCTGGCC ATCTTCGTGCCCATTCACCTGGTGAAGGAGCGGCTCTTCCCTCGGCTGCCACCCGCTTCTGTGGACCATGTGCTG CAGGAGCATCGTGTGGAGCTGCGGCCCACCACGCTGTCGGAGGAGCGGGCACTGCGGGAGCTCGCCCTGCCAGGC TGCACCTCACGCATGCTGAAGTTACTGGCGCTGCGCCAGCTGCCGGACATTTACCCCGACCTTCTCGGCCTGCAG GACAGGTGACTTTGACACTGAGGCTGGAGCTGTGTCCTCCTCAGAGCCCACTGTGGCCAGAGATGAGCCAGAGAG GAAAGCAGAGGCAGCTGCTGGGGAAGAGTCCTGTGGTGCCTCCCCTACCCTGCTACCAGTGCCAGCCCACCTGG CCCCACACTGAAGGCCCGCTTCCGCAGTCTGCTGGAGACCGCCTGGCTCAATGGCCTGGCTCTGCCCACCTGGGG CCACAAGTCCTCAAGACCAGACCAGCCCTCACCCTGCCCACAGCTGCTGGACAGCCAGAGCCATCACCTG**TAG**CA $\tt CTGGTTGCCAGTGCTGTGTATAGCAGTCACTCTCCACCCTTCCCTTCTGCCCAGCTGCCCGGGGCCAC$ GAGTGGATGCTGGGCTGTGCCTCCCCTGGAGGGGTTCCATCTCTGACCCTGTGGCCCATTCAGGGTGGGCT GAAGAGCCCCTGAGCTTTTAACGTGAGGGTCTTTATTGGATAGGACTACTCCCTATTTCTTGCCTAGAGAACACA CATGGGCTTTGGAGCCCGACAGACCTGGGCTTGAATCCCGGCTCGTGTTCTTGCTGCAGGACCTGGGCAAGAAAC ATGCTGCCTGATACATACCCTGTCACCATTTGGTCTCTGCTTCCTCTGGGACAGGGCCTAGAATTGGAGGCAG AGAACCTTCCTATAGAAAGTCTTCGTGTGTCCTAGGACTTGGCTATCGTAGAGTGGTACCTTAGGCAGTGGATGT GACTCACACTCTCAGGAGTCACCCCCCAGCATTTGGGGTTGGGTTGGCCCTACTCCAGCCTGGAGCTCCCTGAGG GAGCCTGCACTCCCTGCTCCCAATCCCCGCTACTGGTGCAGGGATGCAGCCTGGAGCTGGCGTCCTTGTTCTGGG $\tt CCTGCTGCTGCCGCCACCCCAGGAGGCCCCAGGCCTGTCCTGAATTGACATCAGTGCTTCCCTGAACTGCCTCCC$ CCACCCTGGCATTATCCCAGGAAACTTATGTTTTCTAGAAGCTAAGCAGCTGCTGGGACTCAGGGACTGGTGCA GGTAGGCTGAGTGGCAGCTCAGTCCTAGAAGGTCTCTGAAGATCTGGACTGAGGACCCTGCTACTCCCCAAGCCA GAGCCCATCAGCCAGGCCTGTGAGCCACCTGCCTGTGGAGTGCTGAGCTCAACCAAAGGCTGGCAAGCTCTG GGCCTCATTTAAGGGATTCTGATGAGCCGATGGGCCCTGGAGGCAGCCCATTAAAGCATCTGGCTCGTTTTTGGA AAAAAAAAAAAAAA

1016/2825 FIGURE 898

 ${\tt MGQPQWPPELCEQVTGDFDTEAGAVSSSEPTVARDEPESLALAQKSPAPKVRKPGRKPPTPGPEKAEAAAGEESCGASPTPATSASPPGPTLKARFRSLLETAWLNGLALPTWGHKSSRPDQPSPCPQLLDSQSHHL}$

1017/2825 FIGURE 899

1018/2825 FIGURE 900

 ${\tt MGPGPRLLLPLVLCVGLGALVFSSGAEGFRKRGPSVTAKVFFDVRIGDKDVGRIVIGLFGKVVPKTVENFVALAT}$ ${\tt GEKGYGYKGSKFHRVIKDFMIQGGDITTGDGTGGVSIYGETFPDENFKLKHYGIGWVSMANAGPDTNGSQFFITL}$ ${\tt TKPTWLDGKHVVFGKVIDGMTVVHSIELQATDGHDRPLTNCSIINSGKIDVKTPFVVEIADW}$

1019/2825 FIGURE 901

1020/2825 FIGURE 902

 ${\tt MGPGPRLLPLVLCVGLGALVFSSGAEGFRKRGPSVTAKVFFDVRIGDKDVGRIVIGLFGKVVPKTVENFVALAT} \\ {\tt GEKGYGYKGSKFHRVIKDFMIQGGDITTGDGTGGVSIYGETFPDENFKLKHYGIGWVSMANAGPDTNGSQFFITL} \\ {\tt TKPTWLDGKHVVFGKVIDGMTVVHSIELQATDGHDRPLTNCSIINSGKIDVKTPFVVEIADW} \\$

1021/2825 FIGURE 903A

CCACGCGTCCGCCCGGCGCTAAGCCGGGAGACAAGATCCGATCTACTGAGAGCTGCGAGGTACTGCCCTGGATGT GGTGGAGCTGGGCTGGGACCTCCGGGCTGCTGGTCTGTGAGCCTGGCTTTGAAAATTGGTTACTGGGTCTCCTCT CAAGAAATCTCAAGATATTAAAGTTACTGCTGTTAAAAAAGTGCACGAGCATAGCTGTGTCTCGGAAATGGTTGGC GGAAGGTGCAATTTCTCAAGTCGCCTGTTGTTTACATGATGATGATTATGTTGCTGTAGCTACCAGTCAAGGTCT TGTGGTTGTTTGGGAATTAAATCAAGAGCGTCGTGGGAAACCGGAACAAACTGTATGTGTCTTCAGAACACAAAGG CCGAAGAGTCACAGCTCTCTGCTGGGATACAGCTATTCTTAGAGTTTTTGTAGGTGATCATGCTGGGAAGGTTTC TGCTATCAAACTCAATACTTCTAAACAAGCAAAGGCAGCTGCTGCTTTTGTGATGTTTCCTGTTCAGACAATCAC CTTGTGTGACACTGAGAGAAAAGTTTTGGAAAATTGGAAACAAGGAAAGAGATGGAGAATATGGAGCTTGTTT CTTTCCTGGAAGATGTTCTGGGGGCCAGCAACCTCTGATATATTGTGCTCGCCCAGGCTCTAGGATGTGGGAAGT GAACTTTGATGGAGAAGTTATAAGTACACATCAGTTCAAGAAACTCCTCTCGTTGCCACCTCTCCCTGTGATTAC TCTAAGATCAGAACCTCAGTATGATCATACAGCTGGATCCTCCCAGTCTTTGTCTTTCCCCAAACTCTTACATCT TAGTGAGCATTGTGTGCTGACTTGGACAGAAAGAGGGAATTTATATTTTCATTCCTCAGAATGTTCAAGTTCTTCT AGTCTCACATCTCTCCCTGATATCTGTGGAGCGCTGTGTGGAACGCCTGCTAAGAAGAGGCCTATGGAACTTGGC TGCTCGTACATGCTGTCTTTTCCAAAATTCTGTCATTGCCAGCAGAGCAAGAAAAACTTTGACTGCAGATAAATT GGAGCATTTGAAATCTCAGCTGGACCATGGCACCTACAATGATCTAATTTCTCAACTGGAAGAATTGATCTTAAA ATTTGAACCTTTGGATTCAGCTTGTAGCAGTAGAAGAAGCTCCATTTCATCACATGAAAGTTTCAGCATCTTGGA AACCCTCTCAGAAGATGAGAGATTTAAAGAATTCACCTCACAGCAGGAAGAGGACCTGCCAGATCAGTGTTGTGG CTCACACGGAAATGAAGACAATGTTTCTCATGCTCCAGTGATGTTTGAGACAGATAAGAATGAAACTTTTCTCCC CTTTGTGCGTAAAACTACTGAGAAGATTGGCACCCTTCACACGAGCCCTGATCTGAAAGTGAGACCAGAGCTCAG GGGTGATGAGCAATCATGTGAAGAGGATGTGAGTTCAGATACCTGCCCAAAGGAGGAAGACACTGAGGAGGAAAA AGAGGTAACTAGTCCACCTCCAGAAGAAGACAGGTTCCAGGAGCTTAAAGTAGCAACAGCAGAAGCAATGACCAA GCTACAGGACCCTCTGGTTTTATTTGAATCCGAGTCTCTGAGAATGGTTTTACAGGAGTGGCTTTCACATTTAGA AAAAACATTTGCCATGAAGGACTTTTCAGGTGTTTCAGATACTGACAACTCATCCATGAAATTGAACCAGGATGT TGAAGAATCTGTTGATAAAACAGCATGTGAATGTGTAAGGAGTCCAAGGGAGTCTTTGGATGACCTGTTTCAAAT ATGTTCTCCATGCGCCATTGCAAGTGGTCTTCGGAACGACCTGGCTGAATTGACAACATTATGTTTGGAGTTGAA TGTATTGAATTCTAAGATCAAAAGCACCAGTGGACATGTGGACCACACTTTGCAACAGTACTCTCCTGAAATTCT GGCTTGCCAGTTCCTGAAGAAGTACTTTTTCTCCTGAACTTGAAAAGAGCGAAGGAGAGTATCAAGCTTAGTTA GGAGATGGAAAAAGGAGATCTACCAACAAGGTTAAAGTTACTAGATGACGAGGTTCCTTTTGATAGTCCGTTGTT GGTTGTTTATGCTACCCGGTTGTATGAAAAGTTTGGGGAGTCTGCTCTTCGATCCTTAATCAAGTTCTTTCCATC CATTTTGCCATCGGATATCATACAACTTTGTCATCATCATCCTGCTGAGTTTTTGGCCTATTTAGACAGTCTGGT GAAATCAAGGCCTGAAGATCAGCGGTCATCTTTTCTTGAGTCCCTTCTGCAACCAGAGTCTTTAAGGTTGGATTG GCTGCTTTTGGCAGTGTCCCTTGATGCTCCACCAAGCACCAGCACAATGGATGATGAAGGTTATCCCAGGCCTCA TTCACACTTGCTTTCCTGGGGTTACAGTCAGCTGATCCTTCATCTAATTAAACTTCCTGCAGATTTTATAACCAA AGAGAAAATGACAGACATCTGCAGGTCTTGTGGTTTCTGGCCTGGATATCTAATTCTCTGTTTGGAGCTGGAGAG AAGAAGAGGGCCTTCACCAATATTGTGTATCTGAATGATATGAGCCTGATGAAAGGGGACAATGGTTGGATCCC AGAGACCGTGGAGGAATGGAAGCTTCTCCTTCATCTCATACAGAGCAAGAGCACGAGGCCAGCCCCCAGGAGTC ACTAAATGGGAGCCTCAGTGATGGGCCTTCCCCCATCAATGTGGAGAATGTGGCACTTCTGTTAGCTAAGGCCAT GGGCCCAGATCGGGCTTGGTCACTGCTACAGGAATGTGGTCTGGCCCTTGAGTTGTCAGAGAAGTTTACCAGAAC CTGCGATATCCTGAGGATTGCTGAGAAAAGGCAGAGGGCCTTGATACAAAGCATGCTTGAAAAATGCGATCGGTT TCCTGAACCTTCTGAACGCATTTGTTATTGAAGGAAAGACACCACCCCCAAATCCTGCCATCTTATTGGGGGCTAC

FIGURE 903B

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1023/2825 FIGURE 904

MYVSSEHKGRRVTALCWDTAILRVFVGDHAGKVSAIKLNTSKQAKAAAAFVMFPVQTITTVDSCVVQLDYLDGRL
LISSLTRSFLCDTEREKFWKIGNKERDGEYGACFFPGRCSGGQQPLIYCARPGSRMWEVNFDGEVISTHQFKKLL
SLPPLPVITLRSEPQYDHTAGSSQSLSFPKLLHLSEHCVLTWTERGIYIFIPQNVQVLLWSEVKDIQDVAVCRNE
LFCLHLNGKVSHLSLISVERCVERLLRRGLWNLAARTCCLFQNSVIASRARKTLTADKLEHLKSQLDHGTYNDLI
SQMEELILKFEPLDSACSSRRSSISSHESFSILDSGIYRIISSRRGSQSDEDSCSLHSQTLSEDERFKEFTSQQE
EDLPDQCCGSHGNEDNVSHAPVMFETDKNETFLPFGIPLPFRSPSPLVSLQAVKESVSSFVRKTTEKIGTLHTSP
DLKVRPELRGDEQSCEEDVSSDTCPKEEDTEEEKEVTSPPPEEDRFQELKVATAEAMTKLQDPLVLFESESLRMV
LQEWLSHLEKTFAMKDFSGVSDTDNSSMKLNQDVLLVNESKKGILDEDNEKEKRDSLGNEESVDKTACECVRSPR
ESLDDLFQICSPCAIASGLRNDLAELTTLCLELNVLNSKIKSTSGHVDHTLQQYSPEILACQFLKKYFFLLNLKR
AKESIKLSYSNSPSVWDTFIEGLKEMASSNPVYMEMEKGDLPTRLKLLDDEVPFDSPLLVVYATRLYEKFGESAL
RSLIKFFPSILPSDIIQLCHHHPAEFLAYLDSLVKSRPEDQRSSFLESLLQPESLRLDWLLLAVSLDAPPSTSTM
DDEGYPRPHSHLLSWGYSQLILHLIKLPADFITKEKMTDICRSCGFWPGYLILCLELERRREAFTNIVYLNDMSL
MEGDNGWIPETVEEWKLLLHLIQSKSTRPAPQESLNGSLSDGPSPINVENVALLLAKAMGPDRAWSLLQECGLAL
ELSEKFTRTCDILRIAEKRQRALIQSMLEKCDRFLWSQQA

1024/2825 FIGURE 905

 $\texttt{ACCAAGCAAAGCC} \underline{\textbf{ATG}} \texttt{ATATTTCCATGGAAATGTCAGAGCACCCAGAGGGACTTATGGAACATCTTCAAGTTGTG}$ GGGGTGGACAATGCTCTGTTGTGATTTCCTGGCACATCATGGAACCGACTGCTGGACTTACCATTATTCTGAAAA ACCCATGAACTGGCAAAGGGCTAGAAGATTCTGCCGAGACAATTACACAGATTTAGTTGCCATACAAAACAAGGC GGAAATTGAGTATCTGGAGAAGACTCTGCCTTTCAGTCGTTCTTACTACTGGATAGGAATCCGGAAGATAGGAGG AATATGGACGTGGGTGGGAACCAACAATCTCTTACTGAAGAAGCAGAAACTGGGGGAGATGGTGAGCCCAACAA CAAGAAGAACAAGGAGGACTGCGTGGAGATCTATATCAAGAGAAACAAAGATGCAGGCAAATGGAACGATGACGC CTGCCACAAACTAAAGGCAGCCCTCTGTTACACAGCTTCTTGCCAGCCCTGGTCATGCAGTGGCCATGGAGAATG TGTAGAAATCATCAATAATTACACCTGCAACTGTGATGTGGGGTACTATGGGCCCCAGTGTCAGTTTGTGATTCA GTGTGAGCCTTTGGAGGCCCCAGAGCTGGGTACCATGGACTGTACTCACCCTTTGGGAAACTTCAGCTTCAGCTC ACAGTGTGCCTTCAGCTGCTCTGAAGGAACAAACTTAACTGGGATTGAAGAAACCACCTGTGGACCATTTGGAAA CTGGTCATCTCCAGAACCAACCTGTCAAGTGATTCAGTGTGAGCCTCTATCAGCACCAGATTTGGGGATCATGAA CTGTAGCCATCCCCTGGCCAGCTTCAGCTTTACCTCTGCATGTACCTTCATCTGCTCAGAAGGAACTGAGTTAAT TGGGAAGAAAACCATTTGTGAATCATCTGGAATCTGGTCAAATCCTAGTCCAATATGTCAAAAATTGGACAA AAGTTTCTCAATGATTAAGGAGGGTGATTATAACCCCCTCTTCATTCCAGTGGCAGTCATGGTTACTGCATTCTC ATAT**TAA**ATCGCCCTTGGTGAAAGAAATTCTTGGAATACTAAAAATCATGAGATCCTTTAAATCCTTCCATGAA ACGTTTTGTGTGGGGCACCTCCTACGTCAAACATGAAGTGTGTTTCCTTCAGTGCATCTGGGAAGATTTCTACC TGACCAACAGTTCCTTCAGCTTCCATTTCGCCCCTCATTTATCCCTCAACCCCCAGCCCACAGGTGTTTATACAG CTCAGCTTTTTGTCTTTTCTGAGGAGAAACAAATAAGACCATAAAGGGAAAGGATTCATGTGGAATATAAAGATG GAAGTGCAAATTTGATACATATGTGAATATGGACTCAGTTTTCTTGCAGATCAAATTTCACGTCGTCTTCTGTAT GGGTCCTGCTCAAGTTGAAAGAGTCCTATTTGCACTGTAGCCTCGCCGTCTGTGAATTGGACCATCCTATTTAAC TGGCTTCAGCCTCCCCACCTTCTTCAGCCACCTCTCTTTTTCAGTTGGCTGACTTCCACACCTAGCATCTCATGA GTGCCAAGCAAAAGGAGAAGAGAGAAATAGCCTGCGCTGTTTTTTTAGTTTTGGGGGGTTTTGCTGTTTCCTTTTA TGAGACCCATTCCTATTCTTATAGTCAATGTTTCTTTTATCACGATATTATTAGTAAGAAAACATCACTGAAAT GCTAGCTGCAAGTGACATCTCTTTGATGTCATATGGAAGAGTTAAAACAGGTGGAGAAATTCCTTGATTCACAAT GAAATGCTCTCCTTTCCCCTGCCCCCAGACCTTTTATCCACTTACCTAGATTCTACATATTCTTTAAATTTCATC TCAGGCCTCCCTCAACCCCACCACTTCTTTATAACTAGTCCTTTACTAATCCAACCCATGATGAGCTCCTCTTC $\tt CTGGCTTCTTACTGAAAGGTTACCCTGTAACATGCAATTTTGCATTTGAATAAAGCCTGCTTTTTAAGTGTTAA$

1025/2825 FIGURE 906

MIFPWKCQSTQRDLWNIFKLWGWTMLCCDFLAHHGTDCWTYHYSEKPMNWQRARRFCRDNYTDLVAIQNKAEIEY LEKTLPFSRSYYWIGIRKIGGIWTWVGTNKSLTEEAENWGDGEPNNKKNKEDCVEIYIKRNKDAGKWNDDACHKL KAALCYTASCQPWSCSGHGECVEIINNYTCNCDVGYYGPQCQFVIQCEPLEAPELGTMDCTHPLGNFSFSSQCAF SCSEGTNLTGIEETTCGPFGNWSSPEPTCQVIQCEPLSAPDLGIMNCSHPLASFSFTSACTFICSEGTELIGKKK TICESSGIWSNPSPICQKLDKSFSMIKEGDYNPLFIPVAVMVTAFSGLAFIIWLARRLKKGKKSKRSMNDPY

1026/2825 FIGURE 907

1027/2825 FIGURE 908

 ${\tt MTSMTQSLREVIKAMTKARNFERVLGKITLVSAAPGKVICEMKVEEEHTNAIGTLHGGLTATLVDNISTMALLCTERGAPGVSVDMNITYMSPAKLGEDIVITAHVLKQGKTLAFTSVDLTNKATGKLIAQGRHTKHLGN}$

1028/2825 FIGURE 909

TCTGCAGAGATGACGGAGCCCAAGTCGGTGTGTCTCGGTGGATGAGGTGGTGTCCAGCAACATGGAGGCCACT GAGACGGACCTGCTGAATGGACATCTGAAAAAAGTAGATAATAACCTCACGGAAGCCCAGCGCTTCTCCTTCTTG AAAGGATACAAGACCCTCCTGAAAGGAATTTCCGGGAAGTTCAATAGTGGTGAGTTGGTGGCCATTATGGGTCCT ATCAACGGCCTGCCCCGGGACCTGCCTGCTTCCGGAAGGTGTCCTGCTACATCATGCAGGATGACATGCTGCTG CCGCATCTCACTGTGCAGGAGGCCATGATGGTGTCGGCACATCTGAAGCTTCAGGAGAAGGATGAAGGCAGAAGG GAAATGGTCAAGGAGATACTGACAGCGCTGGGCTTGCTGTCTTGCGCCAACACGCGGACCGGGAGCCTGTCAGGT GGTCAGCGCAAGCGCCTGGCCATCGCGCTGGAGCTGGTGAACAACCCTCCAGTCATGTTCTTCGATGAGCCCACC ATTTGCACCATCCACCAGCCCAGCGCCAAACTCTTCGAGCTGTTCGACCAGCTTTACGTCCTGAGTCAAGGACAA TGTGTGTACCGGGGAAAAGTCTGCAATCTTGTGCCATATTTGAGGGATTTGGGTCTGAACTGCCCAACCTACCAC AACCCAGCAGATTTTGTCATGGAGGTTGCATCCGGCGAGTACGGTGATCAGAACAGTCGGCTGGTGAGAGCGGTT $\tt CGGGAGGGCATGTGTGACTCAGACCACAAGAGAGACCTCGGGGGTGATGCCGAGGTGAACCCTTTTCTTTGGCAC$ CGCCCCTCTGAAGAGGTAAAGCAGACAAAACGATTAAAGGGGTTGAGAAAGGACTCCTCGTCCATGGAAGGCTGC CACAGCTTCTCTGCCAGCTGCCTCACGCAGTTCTGCATCCTCTTCAAGAGGACCTTCCTCAGCATCATGAGGGAC GGGAACGAAACCAAGAAGGTCTTGAGCAACTCCGGCTTCCTCTTCTTCTCCATGCTGTTCCTCATGTTCGCGGCC ATCGTGTACTGGATGACGTCGCAGCCGTCCGACGCCGTGCGCTTTGTGCTGTTTTGCCGCGCTGGGCACCATGACC TCCCTGGTGGCACAGTCCCTGGGCCTGCTGATCGGAGCCGCCTCCACGTCCCTGCAGGTGGCCACTTTCGTGGGC $\verb|CCAGTGACAGCCATCCCGGTGCTCCTGTTCTCGGGGGTTCTTCGTCAGCTTCGACACCATCCCCACGTACCTACAG| \\$ TGGATGTCCTACATCTCCTATGTCAGGTATGGGTTCGAAGGGGTCATCCTCTCCATCTATGGCTTAGACCGGGAA GATCTGCACTGTGACATCGACGAGACGTGCCACTTCCAGAAGTCGGAGGCCATCCTGCGGGAGCTGGACGTGGAA CTCAGGTACAAAATCCGGGCAGAGAGG**TAA**AACACCTGAATGCCAGGAAACAGGAAGATTAGACACTGTGGCCGA ${\tt ACCGCGTTGGGTTTGTGGGTGTCTCGTGCTCAGCCACTCTGCCCAGCTGGGTTGGATCTTCTCCATTCCCCTT}$ ACAACTGGGGCAGAATTTAAAGCTGCAACACGCTGGTGATGAGAGGGCTTCCTCAGTCCAGTCGCTCCTTAGCAC CAGAGAGTCCGAGCATGGAGCGATTCCATTTTATGACTGTTGTTTTTCACATTTTCATCTTTCTAAGGTGTGTCT AGTACTTGCTGAAGAATGATTCAGGGTAAATCACATACTTTGTTTAGAGAGGCGAGGGGTTTAACCCGAGTCACC CAGCTGGTCTCATACATAGACAGCACTTGTGAAGGATTGAATGCAGGTTCCAGGTGGAGGGAAGACGTGGACACC ATCTCCACTGAGCCATGCAGACATTTTTAAAAGCTATACACAAAATTGTGAGAAGACATTGGCCAACTCTTTCAA AAAAA

1029/2825 FIGURE 910

MAAFSVGTAMNASSYSAEMTEPKSVCVSVDEVVSSNMEATETDLLNGHLKKVDNNLTEAQRFSSLPRRAAVNIEF RDLSYSVPEGPWWRKKGYKTLLKGISGKFNSGELVAIMGPSGAGKSTLMNILAGYRETGMKGAVLINGLPRDLRC FRKVSCYIMQDDMLLPHLTVQEAMMVSAHLKLQEKDEGRREMVKEILTALGLLSCANTRTGSLSGGQRKRLAIAL ELVNNPPVMFFDEPTSGLDSASCFQVVSLMKGLAQGGRSIICTIHQPSAKLFELFDQLYVLSQGQCVYRGKVCNL VPYLRDLGLNCPTYHNPADFVMEVASGEYGDQNSRLVRAVREGMCDSDHKRDLGGDAEVNPFLWHRPSEEVKQTK RLKGLRKDSSSMEGCHSFSASCLTQFCILFKRTFLSIMRDSVLTHLRITSHIGIGLLIGLLYLGIGNETKKVLSN SGFLFFSMLFLMFAALMPTVLTFPLEMGVFLREHLNYWYSLKAYYLAKTMADVPFQIMFPVAYCSIVYWMTSQPS DAVRFVLFAALGTMTSLVAQSLGLLIGAASTSLQVATFVGPVTAIPVLLFSGFFVSFDTIPTYLQWMSYISYVRY GFEGVILSIYGLDREDLHCDIDETCHFQKSEAILRELDVENAKLYLDFIVLGIFFISLRLIAYLVLRYKIRAER

1030/2825 FIGURE 911

TCTGTGGCCGGAGGCTGATCAGTGTTCTAGAACAGATCAGACATTTTGTAATGATGCCTGAAATAAACACTAACC ACCTCGACAAGCAACAGGTTCAACTCCTGGCAGAGATGTGTATCCTTATTGATGAAAATGACAATAAAATTGGAG CTGAGACCAAGAAGAATTGTCACCTGAACGAGAACATTGAGAAAGGATTATTGCATCGAGCTTTTAGTGTCTTCT TATTCAACACCGAAAATAAGCTTCTGCTACAGCAAAGATCAGATGCTAAGATTACCTTTCCAGGTTGTTTTACGA ATACGTGTTGTAGTCATCCATTAAGCAATCCAGCCGAGCTTGAGGAAAGTGACGCCCTTGGAGTGAGGCGAGCAG GAATTCACTACAAAGCTCAGTCTGATGGTATCTGGGGTGAACATGAAATTGATTACATTTTGTTGGTGAGGAAGA ATGTAACTTTGAATCCAGATCCCAATGAGATTAAAAGCTATTGTTATGTGTCAAAGGAAGAACTAAAAGAACTTC TGAAAAAAGCAGCCAGTGGTGAAATTAAGATAACGCCATGGTTTAAAATTATTGCAGCGACTTTTCTCTTTAAAT GGTGGGATAACTTAAATCATTTGAATCAGTTTGTTGACCATGAGAAAATATACAGAATG<u>TGA</u>ATATGTAGGTAAA TGATTACAGAAAAATTTATGTGCTTAACAAACTTAGAATGACTTTTTCCTTTTAAATTTAGTTCTATCATTAATT TAATTTTGTATCACACCCTGGATATGTGTTCTGTTTCTAAGCGACATTTGTGAGAGATTATTGTAAAATGAGA TTTTAGTAGGCCGTGATAAACATTTCCGGAGCACTTGCAGAGGACTTGCTATTTGCCAGGTGCTTTATGTATCAT TTTCATATTAGCAGATTAAACTAGCAGAGCAAGGTTCAAATTAAATGATATGACCCTAATTTGTATCATTCTGAG TTGATTGTGTGGTTTATTCATTCTGAAACATGTTGATACTTACAGTCACCGACTGCTTTTGATAAGTGATATTGA TTAGGTTGAATCTTCTTGTAAATAGTATTTACCAGTTAGCAAAGTCTGTGTTTTCAGAATTACAGTGAGCACAGA GGTGTTCATAAAATGGGAATTGAGTCCCACTCGGTAAGAGTTGCTTAAACTTGACACTGTTGACATTTGGGCTGG ATAAAACCCCTGTGGTGGGGTCTGTGCTGTGCATTGCAGGATGGTGAGCAGCGTCCCTCTCATGTGACACCCACA GTTATGCCGGATGTTGCCAGATGCCCCTAGGGACAGAGTCAACCCCCAACTGAGGACCACTGTCTACAGAGTCAG GAAATATTGTAGGGAGAAAAAATAACAACAACAAGGCCTATATTAATGTTAAATAGAGGAGATTATGGAATGT AAGACAG

1031/2825 FIGURE 912

MMPEINTNHLDKQQVQLLAEMCILIDENDNKIGAETKKNCHLNENIEKGLLHRAFSVFLFNTENKLLLQQRSDAK ITFPGCFTNTCCSHPLSNPAELEESDALGVRRAAQRRLKAELGIPLEEVPPEEINYLTRIHYKAQSDGIWGEHEI DYILLVRKNVTLNPDPNEIKSYCYVSKEELKELLKKAASGEIKITPWFKIIAATFLFKWWDNLNHLNQFVDHEKI YRM

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1032/2825 FIGURE 913

1033/2825 FIGURE 914

NIGTLSLIK

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1034/2825 FIGURE 915A

GCTGCCCCGAGCCTTTCTGGGGAAGAACTCCAGGCGTGCGGACGCAACAGCCGAGAACATTAGGTGTTGTGGACA GGAGCTGGGACCAAGATCTTCGGCCAGCCCCGCATCCTCCCGCATCTTCCAGCACCGTCCCGCACCCTCCGCATC CTTCCCCGGGCCACCACGCTTCCTATGTGACCCGCCTGGGCAACGCCGAACCCAGTCGCGCAGCGCTGCAGTGAA ${ t TTTTCCCCCCAAACTGCAATAAGCCGCCTTCCAAGGCCAAG{ t ATG}{ t TTCATAAATATAAAGAGCATCTTATGGATGT}$ GTTCAACCTTAATAGTAACCCATGCGCTACATAAAGTCAAAGTGGGAAAAAGCCCACCGGTGAGGGGCTCCCTCT CTGGAAAAGTCAGCCTACCTTGTCATTTTTCAACGATGCCTACTTTGCCACCCAGTTACAACACCAGTGAATTTC TCCGCATCAAATGGTCTAAGATTGAAGTGGACAAAAATGGAAAAGATTTGAAAGAGACTACTGTCCTTGTGGCCC AAAATGGAAATATCAAGATTGGTCAGGACTACAAAGGGAGAGTGTCTGTGCCCACACATCCCGAGGCTGTGGGCG ATGCCTCCCTCACTGTGGTCAAGCTGCTGGCAAGTGATGCGGGTCTTTACCGCTGTGACGTCATGTACGGGATTG AAGACACACAAGACACGGTGTCACTGACTGTGGATGGGGTTGTGTTTCACTACAGGGCGGCAACCAGCAGGTACA CACTGAATTTTGAGGCTGCTCAGAAGGCTTGTTTGGACGTTGGGGCAGTCATAGCAACTCCAGAGCAGCTCTTTG CTCCCAGAGTAGGCTGTTATGGAGATAAGATGGGAAAGGCAGGAGTCAGGACTTATGGATTCCGTTCTCCCCAGG AAACTTACGATGTGTATTGTTATGTGGATCATCTGGATGGTGATGTGTTCCACCTCACTGTCCCCAGTAAATTCA CCTTCGAGGAGGCTGCAAAAGAGTGTGAAAACCAGGATGCCAGGCTGGCAACAGTGGGGGAACTCCAGGCGGCAT GGAGGAACGGCTTTGACCAGTGCGATTACGGGTGGCTGTCGGATGCCAGCGTGCGCCACCCTGTGACTGTGGCCA CTGATAGCAGATTTGATGCCTACTGCTTTAAACCTAAAGAGGCTACAACCATCGATTTGAGTATCCTCGCAGAAA CTGCATCACCCAGTTTATCCAAAGAACCACAAATGGTTTCTGATAGAACTACACCAATCATCCCTTTAGTTGATG AATTACCTGTCATTCCAACAGAGTTCCCTCCCGTGGGAAATATTGTCAGTTTTGAACAGAAAGCCACAGTCCAAC CTCAGGCTATCACAGATAGTTTAGCCACCAAATTACCCACACCTACTGGCAGTACCAAGAAGCCCTGGGATATGG ATGACTACTCACCTTCTGCTTCAGGACCTCTTGGAAAGCTAGACATATCAGAAATTAAGGAAGAAGTGCTCCAGA GCACTGTTTCTGAATTGGTAACCACAGGTCACTATGGATTCACCTTGGGAGAAGAGAGATGATGAAGACAGAACAC TTACAGTTGGATCTGATGAGAGCACCTTGATCTTTGACCAAATTCCTGAAGTCATTACGGTGTCAAAGACTTCAG CTGATAATAATGGATCATCCATGGATGACTGGGAAGAGAGACAAACTAGTGGTAGGATAACGGAAGAGTTTCTTG GCAAATATCTGTCTACTACACCTTTTCCATCACAGCATCGTACAGAAATAGAATTGTTTCCTTATTCTGGTGATA AAATATTAGTAGAGGGAATTTCCACAGTTATTTATCCTTCTACAAACAGAAATGACACATAGAAGAAAAGAA CAGAAACACTAATACCAGAGATGAGAACAGATACTTATACAGATGAAATACAAGAAGAGATCACTAAAAGTCCAT TTATGGGAAAAACAGAAGAAGAAGTCTTCTCTGGGATGAAACTCTCTACATCTCTCAGAGCCAATTCATGTTA CAGAGTCTTCTGTGGAAATGACCAAGTCTTTTGATTTCCCAACATTGATAACAAAGTTAAGTGCAGAGCCAACAG AAGTAAGAGATATGGAGGAAGACTTTACAGCAACTCCAGGTACTACAAAATATGATGAAAATATTACAACAGTGC TTTTGGCCCATGGTACTTTAAGTGTTGAAGCAGCCACTGTATCAAAATGGTCATGGGATGAAGATAATACAACAT CCAAGCCTTTAGAGTCTACAGAACCTTCAGCCTCTTCAAAATTGCCCCCTGCCTTACTCACAACTGTGGGGATGA ATGGAAAGGATAAAGACATCCCAAGTTTCACTGAAGATGGAGCAGATGAATTTACTCTTATTCCAGATAGTACTC AAAAGCAGTTAGAGGAGGTTACTGATGAAGACATAGCAGCCCATGGAAAATTCACAATTAGATTTCAGCCAACTA CATCAACTGGTATTGCAGAAAAGTCAACTTTGAGAGATTCTACAACTGAAGAAAAAGTTCCACCTATCACAAGCA CTGAAGGCCAAGTTTATGCAACCATGGAAGGAAGTGCTTTGGGTGAAGTAGAAGATGTGGACCTCTCTAAGCCAG TACCTTCTGTTCCATCAGAAGATGAAGTTCTAGGTGAACCCTCTCAAGACATACTTGTCATTGATCAGACTCGCC TTGAAGCGACTATTTCTCCAGAAACTATGAGAACAACAAAAATCACAGAGGGAACAACTCAGGAAGAATTCCCTT GGAAAGAACAGACTGCAGAGAAACCAGTTCCTGCTCTCAGTTCTACAGCTTGGACTCCCAAGGAGGCAGTAACAC CACTGGATGAACAAGAGGGCGATGGATCAGCATATACAGTCTCTGAAGATGAATTGTTGACAGGTTCTGAGAGGG TCCCAGTTTTAGAAACAACTCCAGTTGGAAAAATTGATCACAGTGTGTCTTATCCACCAGGTGCTGTAACTGAGC